GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 12, 2005, 22:56:08 ; Search time 164.45 Seconds (without alignments) 45.484 Million cell updates/sec

US-09-932-322-1 44

Title: Perfect score: Sequence: XXXCXPXTGCXXX 13

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

1825181 seqs, 575374646 residues

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Database :

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

. 31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	H	10	9	8	7	თ	ທ	4	ω	N	1	Regult No.
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Q8ECT9	AAR30165	Q6VMH4	Q93KV9	Q8T4I5	Q9WU17	Q9F126	BAD15642 ,	Q6Z8K1	Q926M4	Q92U25	Q80TF6	Q9LEZ5	Q7XV21	AA059515	Q6Y625	Q869K4	Q8LJ36	AAR98366	Q6TVH9	Q9SUV3	Q7U7V9	PA2X_NOTSC	Q726J7	Q9AQP5	Q95KD2	Q9N018	Q7M4J3	Q8KJ72	Q86AK7	Q98AR9	ID
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Aas62716 yersinia		AAS62716	N	151	68.2	30	Ü
Q8zd76 yersinia pe		Q8ZD76	N	151	68.2	30	4
Q6d216 erwinia car	•	Q6D216	N	150	68.2	30	ັພ
Q9liw4 oryza sativ		Q9LIW4	N	133	68.2	30	'n
P08825 bombyx mori		CHA2_BOMMO	_	132	68.2	30	Ξ
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Q8mq78 caenorhabdi		QBMQ78	N	114	68.2	30	8
P08929 bombyx mori		CHA3_BOMMO	.,	100	68.2	30	7
P77130 escherichia		P77130	N	95	68.2	30	5
P90569 plasmodium		P90569	N	78	68.2	30	5
Q7n375 photorhabdu		Q7N375	N	66	68.2	30	4
Q22886 caenorhabdi		Q22886	N	615	70.5	31	ũ
Q91zr8 arabidopsis		Q9LZR8	N	482	70.5	31	2

ALIGNMENTS

% # Q	SO S	R	R R R R R R R R R R R R R R R R R R R	RESULT DE CONTROL OCCUPANTO DE
Query Match 81.8%; Score 36; DB 2; Length 192; Best Local Similarity 71.4%; Pred. No. 27; Matches 5; Conservative 0; Mismatches 2; Indels 0; Gaps 0	"Complete genome structure of the nitrogen-fixing symbiotic bacterium Mesorhizobium loti."; DNA Res. 7:331-338(2000). EMBL; AP003007; BAB52253.1; HSSP; QSZPY9; IFR8. GO; GO:0016491; F:oxidoreductase activity; IEA. GO; GO:001652; P:metabolism; IEA. InterPro; IPR00219; AbH_short. InterPro; IPR00219; AbH_short. Pfam; PF00106; adh short; 1. PRINTS; PR00091; GDHRDH. Complete proteome. SEQUENCE 192 AA; 20183 MW; FD2F660D156037BC CRC64;	UENCE FROM N.A. AIN=MARP303099; LINE=21082930; PubMed=11214968; LINE=21082930; PubMed=11214968; eko T., Nakamura Y., Sato S., Asamizu E., Kato T., Sasamc anabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura I anabe A., Idyokawa C., Kohara M., Matsumoto M., Matsuno A., Nizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto euchi C., Yamada M., Tabata S.;	[1] SEQUENCE PROM N.A. STRAIN=MAFF303099; MEDLINE=21082936; PubMed=1'214974; MEDLINE=21082936; PubMed=1'214974; MEDLINE=21082936; PubMed=1'214974; MEDLINE=21082936; PubMed=1'214974; MATANABE A., Idesawa K., Ishikawa A., Kawashima K., Kimura T., Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T., Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno A., Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto M., Takeuchi C., Yamada M., Tabata S.; Takeuchi C., Yamada M., Tabata S.; Takeuchi C., Yamada M., Tabata S.; Tomplete genome structure of the nitrogen-fixing symbiotic bacterium Mesorhizobium loti (supplement)."; DNA Res. 7:381-406 (2000).	RESULT 1 (998AF) (POPARE) (POPARE
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Q86AK7;
Q86AK7;
Q86AK7;
Q1-JUN-2003 (TrEMBLrel. 24, Created)
Q1-JUN-2003 (TrEMBLrel. 24, Last sequence update)
Q1-JUN-2004 (TrEMBLrel. 26, Last annotation update)
Q1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
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01-OCT-2002
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Submitted (MAR-2003) to the EMBL/GenBank/DDBJ

EMBL; AC116984; AAO51380.1; -.

HSSP; P01382; INTN.

InterPro; IPR0022172; LDL receptor A.

InterPro; IPR001673; S mold repeat.

Pfam; P700526; Dicty_CTDC; 13.

PRINTS; PR00261; LDLRECEPTOR.
                                                                                                                                                   MEDIINE-2199272; PubMed-12003951; Sullivan J.T., Trzebiatowski J.R., Cruickshank R.W., Gouzy J., Brown S.D., Elliot R.M., Fleetwood D.J., McCallum N.G., Rossbach U Stuart G.S., Weaver J.E., Webby R.J., de Bruijn F.J., Ronson C.W.; "Comparative sequence analysis of the symbiosis island of Mesorhizobium loti strain R7A."; J. Bacteriol. 184;3086-3095(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rhizobium loti (Mesorhizobium loti).
Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Phyllobacteriaceae; Mesorhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 25, Last annotation update)
PUTATIVE SHORT-CHAIN TYPE DEHYDROGENASE/REDUCTASE P
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Gloeckner G., Eichinger L., Szafranski K., Pachebat J., Dear Lehmann R., Baumgart C., Parra G., April Jr., Guigo R., Kump Tunggal B., Cox E., Quail M.A., Platzer M., Rosenthal A., Noe "Sequence and analysis of chromosome 2 of Dictyostelium disco Nature 418:79-85 (2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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Bukaryota; Mycetozoa; Dictyosteliida; Dictyostelium
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GO; GO:0008152; P:metabolism; IEA.
InterPro; IPR002198; ADH short.
InterPro; IPR002347; Adh short C2.
Pfam; PF00106; Adh short; 2.
PRINTS; PR00081; GDFRDH.
PRINTS; PR00081; GDFRDH.
PRINTS; PR00081; ADH SHORT; 1.
Oxidoreductase.
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01-MAR-2004
01-MAR-2004
01-MAR-2004
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"Direct induction of Dictyostelium prestalk gene expression provides evidence that DIF is a morphogen.";
Cell 49:185-192(1997).
PIR, A27020, A27020.
InterPro; IPR001673; S. mold_repeat.
Pfam; PF00526; Dicty_CTDC; I9.
NON_TER 601 601
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01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
01-OCT-2003 (TrEMBLrel.
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DIF-induced prestalk pd63 protein precursor (Fragments).
Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.
                                                                                                                                                                                                                                                                                  Hypothetical protein.

Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).

Bukaryota, Metazoa; Chordata; Craniata; Verrebrata; Euceleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
TISSUE=Cerebellum cortex;
Osada N., Hida M., Kusuda J., Tanuma R., Iseki
Osada N., Sugano S., Hashimoto K.;
Suzuki Y., Sugano S., Hashimoto K.;
Submitted (JUL-2000) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                           Cercopithecinae; Macaca.
NCBI_TaxID=9541;
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Best Local
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Q9AQP5;

O1-JUN-2001 (TrEMBLrel. 17, C)

O1-JUN-2001 (TrEMBLrel. 17, L)

O1-MAR-2004 (TrEMBLrel. 26, L)
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Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Bukaryota; Metazoa; Chordata; Craniata; Verebrata; Buteleostomi;
Mammalla; Butheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AB046629; BAB03547.1; -. Hypothetical protein. SEQUENCE 148 AA; 15495 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Medulla oblogata;
Osada N., Hida M., Kusuda J., Tanuma R., Iseki K., Hirai M.,
Suzuki Y., Sugano S., Hashimoto K.;
Submitted (JUN-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AB062948; BAB60737.1; -
HYDOThetical protein.
SEQUENCE 148 AA; 15473 MW; C8D2AA301E0C8191 CRC64;
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Q95KD2; (TrEMBLrel. 19,
01-DEC-2001 (TrEMBLrel. 19,
01-OCT-2003 (TrEMBLrel. 25,
                                                                                                                                                                                                                  MEDLINE-97386424; PubMed-9244273; Sato S., Ouchiyama N., Kimura T., Nojiri H., Yamane H., Omori T.; Sato S., Ouchiyama N., Kimura T., Nojiri H., Yamane H., Omori T.; "Cloning of genes involved in carbazole degradation of Beeudomonas strain CA10: nucleotide sequence of genes and characterization of meta-cleavage enzymes and hydrolase."; J. Bacteriol. 179:4841-4849(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pseudomonas resinovorans.
Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9541;
   Sato S., Nam J., Kasuga K., Nojiri H., "Identification and characterization o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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                                                                                                                                              SEQUENCE FROM N.A.
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                                                                       MEDLINE-97386425; PubMed-9244274;
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RA Klausner R.D., Collins F.S., Wagner L. Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan H., Moore T., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Roha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Hopkinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
M. Jones S.J., Marra M.A.;
Randright R.M., Marra M.A.;
Randright R.M., Marra M.A.;
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Nojiri H., Sekiguchi H., Maeda K., Urata M.
Habe H., Omori T.,
"Genetic characterization and evolutionary
                                                                                                                                                                                                                                                                                                  and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein. Homo sapiens (Human).
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J. Bacteriol. 179:4850-4858(1997).
Strausberg R.;
Submitted (JUN-2003) to the
EMBL; BC053646; AAH53646.1;
Hypothetical protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Lung;
MEDLINE=22388257; PubMed=12477932;
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                                                                                        EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 35; DB 2;
Pred. No. 1.2e+02
                                                                                                                                                                                                                                                                                                           99:16899-16903 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               113 AA
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RESULT 9
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Best Local S
Matches 5
                                                                                                                                                                           InterPro; IPR001211; PhospholipaseA2.

Pfam; PF00068; Phospholip A2 1; 1.

PRINTS; PR00389; PHPHLIPASEA2.

PRODOM; PD000303; PhospholipaseA2; 1.

SMART; SM00085; PA2c; 1.

PROSITE; PS00119; PA2 ASP; 1.

PROSITE; PS00119; PA2 HS; 1.
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                                                                                                                                                                                                                                                                            EMBL; X14043; CAA32201.1; -. PIR; 807983; 807983.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Blapidae; Acanthophiinae; Notechis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Notechis scutatus scutatus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2-acylhydrolase)
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01-PEB-1991 (Rel. 17, Last sequence update)
01-PEB-1991 (Rel. 45, Last annotation update)
Probable phospholipase A2 precursor (EC 3.1.1
                                                                                                                                                                   Calcium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Venom gland;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE PROM N.A.
                                                                                                                                                                                                                                                                  HSSP, P00608, 1AE7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BI_TaxID=70142;
                                                                                                                                                                                                                                                                                                                                                                                                                                      mitted (JAN-1989) to the EMBL/GenBank/DDBJ databases. FUNCTION: PA2 catalyzes the calcium-dependent hydrolysis of acyl groups in 3-sn-phosphoglycerides.

CATALYTIC ACTIVITY: Phosphatidylcholine + H(2)O = 1-acylglycerophosphocholine + a carboxylate.

COPACTOR: Binds I calcium ion per subunit (By similarity).

SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                               subfamily.
                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: Belongs to the phospholipase A2 family. Group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   103
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5; Conserv
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                                                                                                                                                                   Hydrolase;
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                                                                                                                                                                   degradation;
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      By similarity.
Calcium (via carbonyl oxygen) (E
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Pred. No. 40;
                                                                                                        Potential.

Probable phospholipase
By similarity.
                                                                                                                                                         Potent
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(See http://www.isb-sib.ch/announce/
                                                                                                                                                                  Multigene
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                                                                                                                                                                   family; Signal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local &
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9SUV3;
Q9SUV3;
01-MAY-2000
                                                                                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein F8B4.80 (Hypothetical protein AT4g32380).
Name=F8B4.80; Synonyms=AT4g32380;
Arabidopsis thallana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids;
eurosids II, Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

MEDLINB=22825697; PubMed=12917641; DOI=10.1038/nature01943;

Palenik B., Brahamsha B., Larimer F.W., Land M.L., Hauser L.,

Chain P., Lamerdin J.E., Regala W., Allen E.E., McCarren J.,

Pauleen I.T., Dufreene A., Partensky F., Webb E.A., Waterbury

"The genome of a motile marine Synechococcus.";

Nature 424:1037-1042(2003).

EMBL, BX569691; CAE07386.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              METAL
                                                                                       Bevan M., Terryn N., Ardiles W., Buysshav
De Clerck R., De Keyser A., Neyt P., Rou:
Villaroel R., Gielen J., Van Montagu M.,
Mayer K.F.X., Lemcke K., Schueller C.;
Submitted (DEC-1998) to the EMBL/GenBank,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical precursor.
OrderedLocusNames=SYNW0871;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     METAL
SEQUENCE FROM N.A.

EU Arabidopsis sequencing project;
Submitted (AUG-1999) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Complete proteome; Hypothetical protein; Signal SIGNAL 1 22 Potential.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synechococcus sp. (strain WH8102).
Bacteria; Cyanobacteria; Chroococcales; Synechococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2003 (TrEMBLrel. 01-OCT-2003 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q7U7V9;
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=84588;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q7U7V9
                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                    NCBI_TaxID=3702;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 5; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 167 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
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16002
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71.4%;
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                                                                                            the EMBL/GenBank/DDBJ
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Last annotation update)
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Pred. No.
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Calcium (By similarity)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OB3AFB83OECF971C CRC64;
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                                                                                                                                       Buysshaert C., Dassev
P., Rouze P., Van Der
tagu M., Hoheisel J.,
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                                                                                              databases
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                                                                                                                                                                                        Dasseville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                  Den
                                                                                                                                            Mewes H.W.,
                                                                                                                                                                  Daele H.,
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RESULT 12
QATVH
QATVH
AC QATVH
AC QATVH
DT 05-JU
RESULT 13
AAR98366
ID AAR98
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Best Local S
Matches 5
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Best Local 9
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EMBL; AL034567; CAA22565.1; -.
EMBL; AL161581; CAB29955.1; -.
EMBL; T05348; T05348.
GO; GO:0004650; P:polygalacturonase activity; IEA.
GO; GO:0005975; P:carbohydrate metabolism; IEA.
InterPro; IPR000743; Glyco_hydro_28.
InterPro; IPR006626; PbHI.
InterPro; IPR01050; Pectin_lyas_like.
Pfam; PF00295; Glyco_hydro_28; 2.
EMBRT; SM00710; PbHI. 3.
EMBRT; SM00710; PbHI. 3.
EMBRT; SM00710; PbHI. 3.
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Cell wall;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
BUYBSHAEFT C., Dasseville R., De Clerck R.,
De Keyser A., Neyt P., Rouze P., Van Den Daele H., Villaroel R.,
De Keyser A., Neyt P., Rouze P., Van Den Daele H., Villaroel R.,
Gielen J., Van Montagu M., Mewes H.W., Lemcke K., Mayer K.F.X.;
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                          "Genomes of the Parapoxviruses Stomtatitis Virus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE recommends of the EMBL/GenBank/DDBJ databases.
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
Submitted (MAR-2000) to family 28 of glycosyl hydrolases.
  AAR98366
                                                                                                                                                                                                                                                                                                                                                                                  J. Virol. 78:168-177(2004).
EMBL; AY386265; AAR98366.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                   J. Virol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Delhon G., Tulman E.R.,
de la Concha-Bermejillo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUL-2004
05-JUL-2004
                                                                                                                                                                                                                                                                                           Hypothetical
SEQUENCE 4
                                                                                                                                                                                                                                                                                                                                   InterPro; IPR007027; Pox_F11.
Pfam; PF04943; Pox_F11; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN-BV-AR02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bovine papular stomatitis virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=14671098;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rock D.L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TaxID=129727;
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                                                                                                                 26
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5; Conserv
                                                                                                                                                                                                        Similarity 71.
5; Conservative
                                                                                                                 CVPRTGC
                                                                                                                                                         CXPXTGC 10
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0710; PbH1; 3.
Glycosidase; Hydrolase; Hypothetical
312 AA; 34095 MW; E2E70A2622F30EE0
                                                                                                                                                                                                                                                                                             465 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TremBLrel. 27, Created)
(TremBLrel. 27, Last seq
(TremBLrel. 27, Last ann
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[larity 71.4%;
Conservative
    PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
                                                                                                                 32
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                                                                                                                                                                                                                                                                                             51404 MW;
                                                                                                                                                                                                                           77.3%;
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A., Lehmkuhl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA stage;
                                                                                                                                                                                                      Score 34; DB
Pred. No. 1.5e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Last sequence update)
Last annotation update)
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Pred. No. le+02;
0; Mismatches
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  PRT;
                                                                                                                                                                                                                                                                                             BD735469CBED1B79 CRC64;
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  465
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                                                                                                                                                                                                                                                   DB 2;
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..5e+02;
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RMEI, Ap00344; BAC07361.1; -.
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Name=P0413G02.18;
Oryza sativa (japonica cultivar-group).
Eukaryota, Viridiplantae; Streptophyta; Embr Spermatophyta; Magnoliophyta; Liliopsida; Po Ehrhartoideae; Oryzeae; Oryza.
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de la Concha-Bermejillo
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                                                                                                                                                                                                                                Baumgart C.;
Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AC115598; AAO53206.1; -.

R HSSP; P01132; 1EGF.
R GO; GO:0005198; F:structural molecule activity; IEA.
R InterPro; IPR000742; EGF_2.
R InterPro; IPR006209; EGF_1ike.
R InterPro; IPR006210; IEGF.
R InterPro; IPR006210; IEGF.
R InterPro; IPR002049; Laminin_EGF.
R InterPro; IPR002049; Laminin_EGF.
R FARTITS; PF000001; EGF7.7.
R PRINTS; PF000011; EGFLAMININ.
R PROSITE; PS00022; EGF_1; UNKNOWN_10.
R PROSITE; PS00026; EGF_3; 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 77.3
Best Local Similarity 71.4
Matches 5; Conservative
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Best Local Similarity
Matches 5; Conserva
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Q869K4;
Q869K4;
Q1-9UN-2003 (TrEMBLrel. 24, Created)
Q1-JUN-2003 (TrEMBLrel. 24, Last sequence update)
Q1-JUN-2004 (TrEMBLrel. 26, Last annotation update)
G1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Similar to Mus musculus (Mouse). Tenascin X.
Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.
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MEDLINE-23092622; PubMed=12097910;

Gloeckner G.; Eichinger L., Szafranski K., Pachebat J., Dear-P.,

Chemann R., Baumgart C., Parra G., April J.F., Guigo R., Kumpf K.,

Tunggal B., Cox E., Quail M.A., Platzer M., Rosenthal A., Noegel A.A.;

"Sequence and analysis of chromosome 2 of Dictyostelium discoideum.";

Nature 418:79-85(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
STRAIN=AX4;
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STRAIN=AX4;
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                                                                               609 CNPFTGC 615
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                                                                                                                                                        77.3%; Score 34; DB 2; Length 998; llarity 71.4%; Pred. No. 3.1e+02; Conservative 0; Mismatches 2; Indels
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Pred. No. 1.9e+02;
0; Mismatches 2; Indels
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 12, 2005, 23:04:08; Search time 31.2 Seconds (without alignments) 40.090 Million cell updates/sec

Title: Perfect score: Sequence: US-09-932-322-1 44

1 XXXCXPXTGCXXX 13

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

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L	Query Match Best Local Matches	submitted to the EMBL Data Library, A;Reference number: S07983 A;Accession: S07983 A;Molecule type: mxNA A;Residues: 1-145 < DUC> A;Cross-references: UNIPROT:P20146; C;Superfamily: phospholipase A2 F;1-27/Domain: signal sequence #staF;28-145/Product: phospholipase A2	RESULT 2 S07983 phospholipase A2 homolog precursor C;Species: Notechis scutatus scutat (;Date: 07-Sep-1990 #sequence_revicc;Accession: S07983 R;Ducancel; F.	й	Query Match Best Local Matches	A;Molecule type: DNA A;Residues: 1-601 <wil> A;Residues: 1-601 <wil> A;Cross-references: UNIPROT:Q7M4J3 C;Genetics: C;Genetics: A;Gene: pDd63 A;Gene: pDd63 F;1-20/Domain: signal sequence #st F;1-20/Domain: signal sequence #st</wil></wil>	88 ce : , 28 se ce : 1
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	Score Pred. 0; Mi	.46; El	10 E		Score Pred. 0; Mi	M4J3 #status prestalk	in presum presion pres
	re 34; DB d. No. 18; Mismatches	January EMBL:X1 Lus pred	common (common 1 07-2		re 36; DB d. No. 23; Mismatches		protein precursor iscoldeum nce_revision 31-Ma li, A.; McRobbie, of Dictyostelium.p , MUID:87187613; P
	; DB . 18; ches	ary, January 1989 1146; EMBL:X14043; NID #status predicted <sig #status="" a2="" predicted<="" td=""><td>nolog precursor - common tiger snake scutatus scutatus (common tiger sna #sequence_revision 07-Sep-1990 #tex</td><td></td><td>DB 23; ches</td><td>predicted pDd63 pro</td><td>rotein precursor - slime moly soideum 2 revision 31-Mar-1989 #text A.; McRobbie, S.; Mahbuban Dictyostelium prestalk gene</td></sig>	nolog precursor - common tiger snake scutatus scutatus (common tiger sna #sequence_revision 07-Sep-1990 #tex		DB 23; ches	predicted pDd63 pro	rotein precursor - slime moly soideum 2 revision 31-Mar-1989 #text A.; McRobbie, S.; Mahbuban Dictyostelium prestalk gene
	2; L 2;		r sna ger s		2; L 2;	oted <sig> protein #status</sig>	
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	0,	1;	mainland tiger snake) nange 09-Jul-2004		0,	5	(Dictyostelium discoideum) hange 09-Jul-2004 H.; Kay, R.R.; Early, A.; xpression by DIF provides
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hypothetical protein [imported] - Sinorhizobium meliloti (strain 1021) magaplasmid C;Species: Sinorhizobium meliloti
C;Species: Sinorhizobium meliloti
C;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 09-Jul-2004
C;Accession: P96006
R;Finan, T.M.; Weidner, S.; Wong, K.; Buhrmester, J.; Chain, P.; Vorholter, F.J.; H Proc. Natl. Acad. Sci. U.S.A. 98, 9889-9894, 2001
A;Title: The complete sequence of the 1,683-kb pSymB megaplasmid from the N2-fixing A;Accession: P96006
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: T50496
R;Bevan, M.; Terryn, N.; Ardiles, W.; Buysshaert, C.; Dasseville, ewes, H.W.; Rudd, S.; Lemcke, K.; Mayer, K.F.X. submitted to the Protein Sequence Database, May 2000
A;Reference number: Z25101
A;Accession: T50496
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A;Reference number: Z15409
A;Accession: T05348
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R;Bevan, M.; Terryn, N.; Ardiles, W.; Buysshaert, C.; Dasseville, R.; De Clerck, ewes, H.W.; Mayer, K.F.X.; Schueller, C.
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 21-Jul-2000 #sequence_revision 21-Jul-2000 #text_change 09-Jul-2004
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A; Introns: 1/3;
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A;Experimental source: cultivar Columbia; BAC clo
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A, Residues: 1-312 <BEV>
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A,Residues: 1-1058 <BEV>
A,Residues: 1-1058 <BEV>
A,Cross-references: UNIPROT:Q9LEZ5; EMBL:AL357612
A,Experimental source: cultivar Columbia; BAC clone
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,8pecies: Arabidopsis thaliana (mouse-ear cress)
,Date: 23-Apr-1999 #sequence_revision 23-Apr-1999
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;Introns: 60/3; 195/3; 222/3; 448/3; 492/3; 526/3; 555/2; 591/2; 616/3; 662/2;
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A;Cross-references: UNIPROT:(92025; GB:AL591985; PIDN:CAC49718.1; PID:g15141205; GSPDB:A;Experimental source: strain 1021, megaplasmid pSymb
A;Experimental source: strain 1021, megaplasmid pSymb
R;Galibert, F:; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barloy-Hubler
pela, D; Chain, P.; Cowie, A; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fisher, R.F.
L; Hyman, R.W.; Jones, T.
Science 293, 668-672, 2001
A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure
hebault, P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, K.; Yeh, K
A;Title: The composite genome of the legume symbiont Sinorhizobium meliloti.
A;Reference number: A96039; MUID:21368234; PMID:11474104
                                                                                                                                                                                                                                                                                                                                            RESULT 7
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A;Experimental source: cultivar Columbia; BAC clo
C;Genetics:
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A;Accession: T48397
A;Status: preliminary
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R;Bevan, M.; Pohl, T.; Weizenegger,
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A; Residues: 1-64 < KUR>
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-643 <BRA>
A;Residues: 1-643 <BRA>
A;Cross-references: UNIPROT:Q22886; EMBL:U64833; PIDN:AAB04822.1; GSPDB:GN00023; CESP:B
A;Experimental source: strain Bristol N2; clone B0507
                                                                                                                    submitted to the EMBL Data Library, July 1996 A;Description: The sequence of C. elegans cosm A;Reference number: Z20039 A;Accession: T25473
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A; Note: F17C15.120
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                                                                                                                                                                                                                                                                hypothetical protein B0507.1 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #t
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A; Residues: 1-482 <BEV>
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                                                                                                                                                                                                                       R;Bradshaw, H
                                                                                                                                                                                                                                           ;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Gene: SMb21688
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Best Local (
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nes 4; Conservative
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4; Conserv
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57.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 31; DB 2;
Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              r, T.; Bancroft, I.; Mewes, H.W.; Rudd, S.; Database, March 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                       15-Oct-1999 #text_change 09-Jul-2004
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sigma B factor regulatory protein rseC [imported] - Yersinia pestis (strain C;Bpecies: Yersinia pestis
C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004
C;Accession: AB0331
R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.: Px
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Spoerel, N; Nguyen, H.T.; Kafatos, F.C.
J. Mol. Biol. 190, 23-35, 1986
A;Title: Gene regulation and evolution in the chorion locus
A;Reference number: A92929; MUID:87060979; PMID:3023635
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C;Species: Bombyx mori (silkworm)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
C;Accession: B24255
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A;Molecule type: DNA
A;Residues: 1-129 <8PO>
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C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_cha
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_cha
C;Datcossion: A24255
R;Spoerel, N.; Nguyen, H.T.; Kafatos, F.C.
J. Mol. Biol. 190, 23-35, 1986
J. Mol. Biol. 190, 23-35, 1986
A;Title: Gene regulation and evolution in the chorion locus
A;Reference number: A92929; MUID:87060979; PMID:3023635
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C;Species: Bombyx mori (silkworm)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
C;Accession: A24255
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A; Introns: 59/3; 133/1; 464/3; 586/1
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A; Residues: 1-132 <SPO>
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Best Local S
Matches 4
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Best Local
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57.1%;
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Pred. No. 90;
0; Mismatches
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Pred. No. 2e+0
1; Mismatches
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C:Genetics:
A:Gene: rseC
C:Superfamily: Escherichia coli sigma-E factor regulatory protein
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S04926
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A; Map position: 6
C; Superfamily: Neurosp
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A; Molecule type: DNA
A; Residues: 1-199 < SCH>
A; Residues: 1-199 < SCH>
A; Cross-references: EMBL: AL356173; GSPDB: GN00116; NCSP: B14D6.380-
A; Cross-references: BAC clone B14D6; strain OR74A
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A;Gene: win1
A;Introns: 143/3
A;Introns: 143/3
C;Superfamily: hevein precursor; barwin homology; heve:
C;Superfamily: hevein precursor; barwin homology; heve:
F;1-25/Domain: signal sequence #status predicted <SIG>
F;26-200/Product: wound-induced protein 1 #status prediction:
F;26-69/Domain: hevein chitin-binding domain homology.
                                                                                                                                                                                                                                               R;Stanford, A.; Bevan, M.; Northcote, D.

Nol. Gen. Genet. 215, 200-208, Mills a family of novel wound-induced A;Title: Differential expression within a family of novel wound-induced A;Reference number: S04926; MUID:89218921; PMID:2710099
A;Accession: S04926
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submitted to the Protein S
A;Reference number: Z25022
A;Accession: T49486
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A; Residues: 1-151 < KUR>
                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-200 <STA>
                                                                                                                                                                                                                                                                                                                                                                           wound-induced protein 1 precursor - potato
C;Species: Solamum tuberosum (potato)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 09-Jul-2004
C;Accession: S04926
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                                                                                                                                                        C; Genetics:
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57.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 30; DB 2;
Pred. No. 1e+02;
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S.; Barre
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RESULT 15
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C;Species: Nicotiana tabacum (common tobacco)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004
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A,Exporimental source: cultivar Samsun NN
C,Superfamily: hevein precursor; barwin homology; hevein chitin-binding domain homology
P,23-66/Domain: hevein chitin-binding domain homology <HCB>
                                                                                                                                                                                                                                                                                                                                                                                                                    Plant Physiol. 104, 109-118, 1994
A;Title: A novel pathogen- and wound-inducible tobacco
A;Reference number: Z15209; MUID:94159785; PMID:8115541
A;Accession: T04098
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R,Harris, N., Taylor, J.E.; Roberts, J.A.
J. Exp. Bot. 48, 1223-1227, 1997
A,Tille: Characterization and expression of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wound-induced protein (clone TAB7) - tomato (fragment)
C;Bpecies: Lycoporsicon esculentum (tomato)
C;Brecies: Lycoporsicon esculentum (tomato)
C;Date: 14-May.1999 #sequence_revision 14-May-1999 #text_change 09-Jul-2004
C;Accession: T07729
                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-211 < PON>
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A;Accession: T07729
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  wound-induced
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A;Molecule type: mRNA
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Ponstein, A.S., Bres-Vloemans, S.A.; Sela-Buurlage, M.B.; Elzen, P.J.; Melchers,
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C;Superfamily: hevein precursor; barwin homology; hevein chitin-binding F;1-25/Domain: signal sequence #status predicted <SIG> F;26-211/Product: wound-induced protein 2 #status predicted <MAT> F;26-69/Domain: hevein chitin-binding domain homology <HCB> F;77-198/Domain: barwin homology <BAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mol. Gen. Genet. 215, 200-208, 1989
A; Title: Differential expression within a family of novel wound-induced A; Reference number: S04926; MUID:89218921; PMID:2710099
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C;Date: 28-Feb-1990 #sequence_revision 2/
C;Accession: S04927
C;Accession: S04927
R;Stanford, A.; Bevan, M.; Northcote, D.
Search completed: January 12, 2005, 23:16:03 Job time: 33.2 secs
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A;Residues: 1-211 <STA>
A;Cross-references: UNIPROT:P09762; EMBL:X13497; NID:g21617;
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and is derived by analysis of the total score distribution.
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                     ABG28408
AAU40006
ABM36525
AAG43626
AAG43625
                                                                             ABB93198
ADF05702
                                                                                                  ABJ00545
ABG33861
AAU44932
ABM41451
ADA54961
AAB42027
ABO75329
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                                                                                                                                                                                                                                                                                                                                 SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Search time 146.9 Seconds
(without alignments)
31.746 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002273
                             Aau49413 Propionib
Abm4593 Propionib
Abm74428 Pseudomon
Aay62764 Pseudomon
Aay62764 Pseudomon
Aay62007 Cadherin-
Aay62024 Cadherin-
Aay62224 Cadherin-
Abj00550 B lymphoc
Abj33862 B lymphoc
Abj33861 B lymphoc
Abj40545 Propionib
Adm41451 Propionib
Adm41451 Propionib
Adm54961 Human pro
Aab42027 Human Orr
Aab42027 Human Orr
Aab42027 Human Orr
Aab42027 Human Orr
Aab42027 Bugetriid
Adm615702 Bacterial
Abg28408 Novel hum
Aau40006 Propionib
Abm36525 Propionib
Abm36525 Propionib
Abm36525 Propionib
Abm36525 Propionib
Abm36525 Propionib
Abm36525 Propionib
                                                                                                                                                                                                                                                                                             Description
 Propionib
Arabidops
Arabidops
Arabidops
Protocadh
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26
31	31	31	31	3	31	31	31	<u>3</u> 1	2	31	31	31	31	31	31	31	31	31	31
70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5	70.5
638	602	580	580	580	580	580	580	482	444	310	239	239	238	193	193	183	148	114	90
N	N	N	N	N	N	N	N	v	7	7	σ	σ	σ	7	4	7	7	σ	7
AAY04992	AAY04996	AAY39105	AAY38968	AAW81666	AAW64303	AAW32431	AAW32363	ABB93340	ABO70819	ABO74638	ABP76713	ABP99348	ABP99259	ADF60298	AA000015	ADF58925	ADJ71040	ABP06368	AB081251
Aay04992 Mycobacte	Aay04996 Mycobacte	Aay39105 M. tuberc	Aay38968 M. tuberc	Aaw81666 M. tuberc	Aaw64303 Mycobacte	Aaw32431 Mycobacte	Aaw32363 Mycobacte	Abb93340 Herbicida	Abo70819 Pseudomon	Abo74638 Pseudomon	Abp76713 Streptomy	Abp99348 Orthosomy	Abp99259 Orthosomy	Adf60298 Human con	Aao00015 Human pol	Adf58925 Human pol	Adj71040 Human hea	Abp06368 Human ORF	Abo81251 Pseudomon

ALIGNMENTS

AAU49413 standard; protein; 79 A

AAU49413;

13-FEB-2002 (first entry)

Propionibacterium acnes immunogenic protein #10309

SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.

Propionibacterium acnes.

WO200181581-A2

01-NOV-2001

20-APR-2001; 2001WO-US012865.

21-APR-2000; 2000US-0199047P 02-JUN-2000; 2000US-0208841P 07-JUL-2000; 2000US-0216747P

(CORI-) CORIXA CORP.

Skeiky YAW, Persing DH, L'maisonneuve J, Zhang Y, Mitcham JL, Wang SS, , Jen S, Carter D;

WPI; 2001-616774/71. N-PSDB; AAS59545.

Propionibacterium acnes vaccinating against and treating acne vulgaris. polypeptides and nucleic acids useful for diagnosing infections, especially useful for

Example 1; SEQ ID NO 10608; 1069pp; English.

RESULT 1
AAU49413
AID AAU49413
AIC AAU44
AIX AAU44
AIX AAU4
AIX AA Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory

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ARBSULT 2
ARBMAS 2
ARBMAS 2
ARBMAS ARMA ACC
ARBMA ACC
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Best Local
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The invention relates to an isolated polynucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates to polypeptides encoded by the polynucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynucleotide of the invention; antibodies against polypeptides of the invention; a method for stimulating an immune response specific for a P. acnes polypeptide and an isolated T cell population comprising T cells prepared via this method; a vaccine composition (comprising P. acnes polypeptides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitcham JL, Skeiky YAW, Pe
Zhang Y, Wang S, Jen S, I
Barth B, Vallieve-Douglass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Acne vulgaris; antiseborrhoeic; dermatological; antibacterial; immunostimulant; immune response; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 79 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptide, useful for diagnosing, preventing or for stimulating an immune response specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-2001; 2001US-00978825
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                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 10608; 1481pp; English.
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5, Conserv
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ang S, Jen S, Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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Pred. No. 55;
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Benson DR,
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Jones R, Carte
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                                                                               cc polynucleotides, antibodies, fusion proteins, T cell populations, or cantigen-presenting cells that express the polypoptide); a method and kit cf for detecting or determining the presence or absence of P. acnes in a cc patient; and a method for inhibiting the development of P. acnes in a patient. The P. acnes polypoptides, polynucleotides, antibodies, fusion croteins, T cell populations or antigen-presenting cells that express the polypoptides are useful for diagnosing, preventing or treating acne conjugaris, or for stimulating an immune response specific for a P. acnes for cultaris, or for stimulating an immune response specific for a P. acnes crotein. The polynucleotides can also be used as probes or primers for nucleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response against P. acnes, or for treating acne, can the kit is useful for performing a diagnostic assay. The present contained within the P. acnes polynucleotides of the cinvention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly contained within the P. acnes polynucleotides of the contained within the P. acnes polynucleotides of the printed specification, but was obtained in electronic format directly contained within the P. acnes polynucleotides of the contained within the P. acnes polynucleotides of the printed specification, but was obtained in electronic format directly contained within the P. acnes polynucleotides of the printed specification, but was obtained in electronic format directly contained within the P. acnes polynucleotides of the contained within the P. acnes polynucleotides of the printed specification, but was obtained in electronic format directly contained the present and the contained within the P. acnes polynucleotides of the printed specification.
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                                         Query Match
              4 CXPXTGC 10
                          Similarity 5; Conserv
CSPGTGC 9
                           Conservative
                                   79.5%;
71.4%;
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                            Score 35; DB Pred. No. 55; O; Mismatches
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AXX ARBO7
AXX ARBO7
AXX ARBO7
AXX Bact
XXX Bact
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XXX C22-1
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Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.

Pseudomonas aeruginosa

US6551795-B1

18-FEB-1999;

99US-00252991

18-FEB-1998; 27-JUL-1998; 98US-0074788P 98US-0094190P

(GENO-) GENOME THERAPEUTICS CORP

Rubenfield MJ, Nolling Ç Deloughery Ç Bueh

N-PSDB; 2003-615309/58. ABD07999

Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment o pathological conditions resulting from bacterial infection. Of

Disclosure; SEQ ID NO 23174; 455pp; English

The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant

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                                                                                                                                                                                                                                                                               05-MAY-1998;
06-NOV-1998;
20-JAN-1999;
 The present invention describes cadherin modulating agents (MA) comprising peptides which comprise a nonclassical cadherin cell adhrecognition (CAR) sequence. The MAs can be used for modulating nonclassical cadherin-mediated functions. They can be used for e.g. inhibiting adhesion of nonclassical-cadherin expressing cells in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inhibition; cadherin extracellular domain; cell adhesion; recognition; OB-cadherin; cadherin-5; cadherin-6; cadherin-7; cadherin-8; cadherin-12; cadherin, cadherin-15; T-cadherin; PB-cadherin; cadherin; cadherin; cadherin; cadherin; protocadherin; desmocollin; calcium binding; cancer; tumour; obesity; rheumatoid arthritis; multiple sclerosis; diabetes; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 180 AA;
                                                                                                                   New cadherin modulating agents, used for modulating nonclassical cadherin -mediated functions for treating e.g. cancers, obesity, rheumatoid arthritis, multiple sclerosis, diabetes or a neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY62764 standard; peptide; 7
                                                                                                                                                                                                                                                                                                                                           05-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PB-cadherin cell adhesion recognition cyclic peptide SEQ
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                                                                                                                                                                                                                                                                                                                                                                        11-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DINO
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                                                                                                                                                                                                                                     (ADHE-) ADHEREX TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens
                                                                                                                                                                               2000-038791/03
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                                                                                      72; Page 193; 252pp; English
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
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                                                                                                                                                                                                                                                                   98US-00073040.
98US-00187859.
99US-00234395.
99US-00264516.
                                                                                                                                                                                                                                                                                                                                           99WO-CA000363.
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1. .7
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71.4%;
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                                                                                                                                                                                                          Byers S;
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Pred. No. 1.1e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            adhesion; CAR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID NO:4047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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                                               adhesion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC enhancing delivery of a drug to a tumour in a mammal, treating cancer in Ca mammal, inhibiting metastasis of a cancer in a mammal, inhibiting cancer in a mammal, inhibiting cangiogenesis in a mammal in a mammal, inhibiting cangiogenesis in a mammal apoptosis in a nonclassical cadherince expressing cell, preventing or treating obesity in a mammal, stimulating clood vessel regression in a mammal anhancing drug delivery to the contral nervous system, treating a demyelinating neurological disease, concreasing vasopermeability in a mammal, enhancing adhesion of conclassical cadherin-expressing cells, inhibiting synaptic stability in conclassical cadherin-expressing cells, inhibiting synaptic stability in conclassical cadhering preventing preventing perlia, inhibiting synaptic stability in conclassical reventing preventing neurite outgrowth, facilitating wound healing cor reducing scar tissue, or enhancing adhesion of foreign tissue in a commmal. They can also be used for treating e.g. psoriasis, arthritis, age crelated macular degeneration, multiple sclerosis and diabetes. The products can also be used for detection and diagnosis and in bioreactors. ANY66592 to ANY64572 represent specifically claimed peptides, and cany64573 to ANY64643 and ANZ33183 to ANZ33186 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Loc
Matches
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                                                                                                                                                                                                                                                        05-MAY-1998;
06-NOV-1998;
20-JAN-1999;
08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modulation; nonclassical cadherin mediated cell adhesion; CAR; inhibition; cadherin extracellular domain; cell adhesion recognition; OB-cadherin-5; cadherin-6; cadherin-7; cadherin-8; cadherin-12; cadherin-14; cadherin-15; T-cadherin; PB-cadherin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mammal, enhancing delivery of a drug through the skin of a
                                                                                                                            Blaschuk OW,
                                                                                                                                                                                                                                                                                                                                                                                                                          05-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9957149-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rheumātoid arthritis; multiple sclerosis; diabetes; metastasis; neurological disease; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cadherin related neuronal receptor; LI-cadherin; protocadherin;
desmoglein; desmocollin; calcium binding; cancer; tumour; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cadherin-7 cell adhesion recognition cyclic peptide SEQ ID NO:1375.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY61489;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY61489 standard; peptide; 7
                                                                                                                                                                                               (ADHE-)
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5; Conserv
                                                                                                                                                                                            ADHEREX TECHNOLOGIES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CXPXTGC 10
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                                                                                                                            Gour BJ,
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                                                                                                                                                                                                                                                            98US-00073040.
98US-00187859.
99US-00234395.
99US-00264516.
                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-CA000363
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1. .7
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Pred. No. 1.7e+06;
0; Mismatches 2
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New cadherin modulating agents, used for modulating nonclassical cadherin

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AAX762007
ID AAX76
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XX AAX6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modulation; nonclassical cadherin mediated cell adhesion; CAR; inhibition; cadherin extracellular domain; cell adhesion recognition; OB-cadherin; cadherin-5; cadherin-6; cadherin-7; cadherin-8; cadherin cadherin-14; cadherin-15; T-cadherin; PB-cadherin; cadherin; protocadherin; cadherin related neuronal receptor; LI-cadherin; protocadherin; desmoglein; desmocollin; calcium binding; cancer; tumour; obesity; rheumatoid arthritis; multiple sclerosis; diabetes; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   comprising peptides which comprise a nonclassical recognition (CAR) sequence. The MAs can be used to nonclassical cadherin-mediated functions. They can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes cadherin modulating agents comprising peptides which comprise a nonclassical cadherin
    05-MAY-1998;
06-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cadherin-12 cell adhesion recognition cyclic peptide SEQ ID NO:1799
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY62007 standard; peptide; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 36; Page 172; 252pp; English.
                                                                                                                    05-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neurological
                                                                                                                                                                                                   11-NOV-1999.
                                                                                                                                                                                                                                                                                  WO9957149-A2
                                                                                                                                                                                                                                                                                                                                                             Disulfide-bond
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5, Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
        98US-00073040
98US-00187859
                                                                                                                        99WO-CA000363
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 34; DB 3;
Pred. No. 1.7e+06;
0; Mismatches 2
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cell adhesion
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ANY62224
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ANY62224
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Best Local &
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                                                                                                                                                                                                                        Modulation; nonclassical cadherin mediated cell adhesion; CAR; inhibition; cadherin extracellular domain; cell adhesion recognition; OB-cadherin-1; cadherin-1; cadherin-1; cadherin-1; cadherin-1; cadherin-1; cadherin-1; cadherin-1; cadherin-14; cadherin-15; T-cadherin; PB-cadherin;
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08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                  Cadherin-14 cell adhesion recognition cyclic peptide SEQ ID NO:3971.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY62224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY62224 standard; peptide; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the exemplification of the present invention
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5; Conserv
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99US-00264516.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 34; DB 3;
Pred. No. 1.7e+06;
0; Mismatches 2
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cadherin related neuronal receptor; LI-cadherin; protocadherin;
desmoglein; desmocollin; calcium binding; cancer; tumour; obesity;
rheumatoid arthritis; multiple sclerosis; diabetes; metastasis;

Synthetic. neurological

disease;

cyclic.

Homo

sapiens

Location/Qualifiers

Disulfide-bond

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ARBJOULT 8
ARBJOOSSO
ID ARBJO
XX ARBJO
XX ARBJO
XX OS-9
XX OS-9
XX B ly
XX B ly
XX Alle
KW alle
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                                                                                                                                                                                                                                                                                                                                                                                                                                  CC nonclassical cadherin-mediated functions. They can be used for e.g. CC inhibiting adhesion of nonclassical-cadherin expressing cells in a CC mammal, enhancing delivery of a drug through the skin of a mammal, CC enhancing delivery of a drug to a tumour in a mammal, treating cancer in CC amammal, inhibiting metastasis of a cancer in a mammal, inhibiting metastasis of a cancer in a mammal, inhibiting color color color color cells in a mammal, inducing apoptosis in a nonclassical cadherin-cc expressing cell, preventing or treating obesity in a mammal, stimulating CC blood vessel regression in a mammal, enhancing drug delivery to the CC central nervous system, treating a demyelinating nurrological disease, increasing vasopermeability in a mammal, enhancing adhesion of cc a mammal, or preventing pregnancy in a mammal. They can also be used for ce.g. enhancing scartispus, or enhancing adhesion of foreign tissue in a cc mammal. They can also be used for treating e.g. psoriasis, arthritis, age crelated macular degeneration, multiple sclerosis and diabetes. The crelated macular degeneration, multiple sclerosis and in bioreactors. AAY60592 to AAY64573 and AAZ33183 to AAZ33186 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-MAY-1998;
06-NOV-1998;
20-JAN-1999;
08-MAR-1999;
B lymphocyte stimulator protein binding protein; BLyS; immune disease; allergy; proliferative disease; infectious disease; arteriosclerosis; inflammatory disorder; hypergammaglobulinaemia; blood clotting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New cadherin modulating agents, used for modulating nonclassical c-mediated functions for treating e.g. cancers, obesity, rheumatoid arthritis, multiple sclerosis, diabetes or a neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising peptides which comprise a nonclassical cadherin cel recognition (CAR) sequence. The MAs can be used for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 54, Page 184; 252pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-038791/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blaschuk OW,
                                                                                                           05-SEP-2002
                                                                                                                                                                             ABJ00550 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ADHE-) ADHEREX TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 5; Conserv
                                                                      lymphocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention describes cadherin modulating agents
                                                                                                                                                                                                                                                                                                 CXPXTGC 10
                                                                                                                                                                                                                                                                  CDPKTGC 7
                                                                                                                                                                                                                                                                                                                                                                                                         A
A
                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                    stimulator protein binding peptide #1.
                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-00187859.
99US-00234395.
99US-00264516.
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                                                                                                                                                                           peptide; 7
                                                                                                                                                                                                                                                                                                                                                   77.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Byers S;
                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                  Score 34; DB 3;
Pred. No. 1.7e+06;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                      2
                                                                                                                                                                                                                                                                                                                                                                     Length 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell adhesion
                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                      Gaps
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ischaemia; graft-versus-host disease; neurodegenerative disease; immunosuppressive; nephrotropic; antirheumatic; antiarthritic; neuroprotective; cytostatic; immunostimulant; antitumour; anti-HIV; antiasthmatic; antiallergic; thyromimetic; antianaemic; haemostatic; dermatological; antianflammatory; cardiant; ophthalmological; uropathic; antidiabetic; antithyroid; antidepressant; hepatotropic.
                                                                                                                                                                     Claim
                                                                                                                                                                                                                                           Beltzer
                                                                                                                                                                                                                                                                                    18-AUG-2000; 2000US-0226700P
                                                                                                                                                                                                                                                                                                        17-AUG-2001; 2001WO-US025850
                                                                                                                                                                                                                                                                                                                             28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                  WO200216411-A2
                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                         administering
                                                                                                                                                                                                                                                                (HUMA-)
                                                                                                                                                                                      treatment of various diseases e.g.
inistering B Lymphocyte stimulator b
                                                                                                                                                                                                                       2002-499775/53
                                                                                                                                                                   69; Page 233; 387pp; English
                                                                                                                                                                                                                                           JP,
                                                                                                                                                                                                                                                                HUMAN GENOME SCI
                                                                                                                                                                                                                                           Potter DM,
                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                      /label= Pro, Tyr
                                                                                                                                                                                                                                                                                                                                                                                           label= Phe,
                                                                                                                                                                                                                                          Fleming TL, Rosen CA;
                                                                                                                                                                                        binding polypeptide
                                                                                                                                                                                                   rheumatoid arthritis, comprises
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Query Match
Best Local S
Matches 7
                                                                                                                                                                                                                           amelioration of a disease or disorder associated with: aberrant B lymphocyte stimulator (BLyS), BLyS receptor expression or activity; cells of haematopoietic origin; or proliferative disease; and reducing, inhibiting or stimulating immunoglobulin production, B cell proliferation and graft rejection involving administration of BLyS binding polypeptide. The BLyS binding polypeptides are used in the treatment, prevention or amelioration of diseases such as immune system diseases, proliferative diseases, diseases of cells of hematopoietic origin, graft rejection, allergies, infectious diseases, arteriosclerosis, inflammatory disorders, hypergammaglobulinaemia, blood clotting disorders, ischaemia, and neurodegenerative diseases. The present sequence is a B lymphocyte stimulator protein binding peptide
                                                                                                                                                                                             Sequence
                                                         Similarity 7; Conserv
CXPXTGC 10
                                                                                                                                                                                                AA;
                                                            Conservative
                                                            77.3%; Score 34; DB 5; L-
100.0%; Pred. No. 1.7e+06;
Live 0; Mismatches 0;
                                                                                                                            Length 7;
                                                                   0
                                                                   Gaps
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RESULT 9
ABG33862
ID ABG33862
AC ABG3
XX ABG3
XX
DT 15-J
DT 15-J
DX B Ly
XX
XX
XX
B Ly
XX
XX
XX
XX
XX
SYNO
XX

15-JUL-2002

(first

entry)

BLyS;

Lymphocyte Stimulator protein; B Lymphocyte Stimulator binding LyS; biological fluid; serum; plasma; lymph; blood; urine; spina ynovial fluid; saliva; mucus.

urine; spinal

ABG33862;

ABG33862 standard; peptide;

B Lymphocyte Stimulator (BLyS) binding peptide #436

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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a B Lymphocyte Stimulator (BLyS) binding polypeptide. BLyS binding peptides bind BLyS or BLyS-like proteins reversibly or irreversibly. The binding peptides are used in detection, isolation and/or purification of BLyS in a solution such as water or a buffer solution, as well as any fluid and/or cell obtained from an individual biological fluid, body tissue, body cell, cell line, tissue culture or other source containing BLyS or BLyS-like polypeptides. The biological fluid, sera, plasma, lymph, blood, blood fraction, urine, synovial fluid, spinal fluid, selliva and mucous. Sequences ABG33406-33415, ABG33423-33575, ABG3358-33846, ABG33848-33850 and ABG33852-33862 represent BLyS binding peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New B-lymphocyte stimulator binding polypeptide useful in detecting isolating BLyS or BLyS-like polypeptide comprises a specified amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                             B lymphocyte stimulator protein binding protein; BLyS; immune disease; allergy; proliferative disease; infectious disease; arteriosclerosis; inflammatory disorder; hypergammaglobulinaemia; blood clotting; ischaemia; graft-versus-host disease; neurodegenerative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence.
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Misc-difference
                                                                                                                                                                                                                                                                                                                         05-SEP-2002
                                                                                                                                                                                                                                                                                                                                                      ABJ00545
                                                                                                                                                                                                                                                                                                                                                                                     ABJ00545 standard; peptide; 13
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                             Misc-difference
                                                             Misc-difference
                                                                                                       Unidentified
                                                                                                                                       antidiabetic;
                                                                                                                                                   neuroprotective; cytostatic; immunostimulant; antitumour; anti-HIV; antiasthmatic; antiallergic; thyromimetic; antianaemic; haemostatic; dermatological; antiinflammatory; cardiant; ophthalmological; uropathic;
                                                                                                                                                                                                                                                                                          B lymphocyte
                                                                                                                                                                                                lschaemia; graft-versus-host disease; neurodegenerative disease;
lmmunosuppressive; nephrotropic; antirheumatic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CXPXTGC 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 132; 269pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORP
                                                                                                                                                                                                                                                                                         stimulator protein binding protein consensus peptide #1.
                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                      antithyroid; antidepressant;
                                                                          Location/Qualifiers
            /label= Ala, Glu, Met, Ser, Val
                                          label- Ala, Asn, Lys,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77.3%; Score 34; DB
100.0%; Pred. No. 1.
ive 0; Mismatches
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thes 0;
                                             Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 7;
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                                                                                                                                                                                   anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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acid
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RESULT 11
ABG33861
ID ABG33861
AC ABG33
XX ABG33
XX ABG33
XX ABG33
XX ABG33
XX BBJ9T
DT 15-JT
DT 15-JT
DT BLym
XX BLym
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of haematopoietic origin; or proliferative disease; and reducing, inhibiting or stimulating immunoglobulin production, B cell proliferation and graft rejection involving administration of BLyS binding polypeptide. The BLyS binding polypeptides are used in the treatment, prevention or amelioration of diseases such as immune system diseases, proliferative diseases, diseases of cells of hematopoietic origin, graft rejection, allergies, infectious diseases, arteriosclerosis, inflammatory disorders, hypergammaglobulinaemia, blood clotting disorders, ischaemia, and neurodegenerative diseases. The present sequence is a conserved region of a B lymphocyte stimulator protein binding peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                               B Lymphocyte Stimulator protein; B Lymphocyte Stimulator binding peptide; BLyS; biological fluid; serum; plasma; lymph; blood; urine; spinal fluid; synovial fluid; saliva; mucus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to the treatment, prevention or amelioration of a disease or disorder associated with: aberra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The treatment of various diseases e.g. rheumatoid arthritis, comprises administering B Lymphocyte stimulator binding polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beltzer JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-AUG-2000; 2000US-0226700P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
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                                                                                                                                                                                                      B Lymphocyte Stimulator
                                                                                                                                                                                                                                                        15-JUL-2002
                                                                                                                                                                                                                                                                                                        ABG33861;
                                                                                                                                                                                                                                                                                                                                                       ABG33861 standard; peptide; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 69; Page 230; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200216411-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amelioration of a disease or disorder associated with: aberrant B
lymphocyte stimulator (BLyS), BLyS receptor expression or activity; cells
of haematopoletic origin; or proliferative disease; and reducing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HUMA-) HUMAN GENOME SCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
7; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 CXPXTGC 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CXPXTGC 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /label=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label=
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                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                                      (BLyS) binding peptide #435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 34;
Pred. No.
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BG
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Synthetic

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ARESULT 12
AAU44932
ID AAU44
XX AU44
AX AAV
AX AVA
AX SAPHC
KW UVGIT
KW Infla
KW Infla
KW O200
XX Propi
YX O1-W
YX O1-
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Best Local &
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                                                                                                                                          21-APR-2000;
02-JUN-2000;
07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; EIISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a B Lymphocyte Stimulator (BLyS) binding polypeptide. BLyS binding peptides bind BLyS or BLyS-like proteins reversibly or irreversibly. The binding peptides are used in detection, isolation and/or purification of BLyS in a solution such as water or a buffer solution, as well as any fluid and/or cell obtained from an individual biological fluid, body tissue, body cell, cell line, tissue culture or other source containing BLyS or BLyS-like polypeptides. The biological fluids include sera, plasma, lymph, blood, blood fraction, urine, symovial fluid, spinal fluid, saliva and mucous. Sequences ABG33406-33415, ABG33433-33575, ABG33588-33846, ABG33848-33850 and ABG33852-33862 represent BLyS binding peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                      WO200181581-A2
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     Skeiky YAW, F
L'maisonneuve
                                                                                                                                                                                                                                                          20-APR-2001; 2001WO-US012865.
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                                                                                                                                                                                                                                                                                                                                                                                                                           Propionibacterium acnes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beltzer JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-AUG-2000; 2000US-0226489P
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                                                                                  (CORI-) CORIXA CORP.
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  Persing DH,
e J, Zhang Y,
                                                                                                                                          2000US-0199047P.
2000US-0208841P.
2000US-0216747P.
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Mitcham JL, Wang SS,
, Jen S, Carter D;
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Pred. No.
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RESULT 13
ABM41451
ID 41841
XX ABM41
XX ADM41
XX ACNe
DE Propi
XX ACNe
XW immun
XX W0200
XX Propi
XX W0200
XX 11-00
XX I1-00
XX I1-00
XX II-00
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XX III-00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cc polypeptides. The proteins and their associated DNA sequences are used in cthe treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, cc. pustulosis, hypertosis and osteomyelltis), uveitis and endophthalmitis. CP. acnes is also involved in infections of bone, joints and the central creations associated with acne vulgaris. A method for detecting the creations associated with acne vulgaris. A method for detecting the creating the amount of bound protein in the invention and determining the amount of bound protein in the sample. The cand determining the amount of bound protein in the sample to the proteins of the invention and determining the amount of bound protein in the sample to the proteins. The collection of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and ctherefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by cenzyme linked immunosorben. assay (BLISA). Note: The sequence data for this patent did not form part of the printed specification, but was contact of the printed specification, but was contact of the printed specification, but was contact of the printed specification.
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Best Local
                                                                                     Mitcham JL, Skeiky YAW, Pers
Zhang Y, Wang S, Jen S, Lod
Barth B, Vallieve-Douglass J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Acne vulgaris; antiseborrhoeic; dermatological; immunostimulant; immune response; vaccine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID NO 6127; 1069pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
      N-PSDB; ACF64453
                                 WPI; 2003-381789/36.
                                                                                                                                                                                                                                                                                                                                             11-OCT-2002; 2002WO-US032727.
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                                                                                                                                                                                                                                                                                  15-OCT-2001; 2001US-00978825
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                                                                                                                                                                                                                      (CORI-)
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71.4%;
                                                                                                                          Persing DH,
Lodes MJ,
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Pred. No. 60;
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                                                                                                                          Bhatia A,
Benson DR,
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                                                                                                                          Maisonneuve
Jones R, Ca
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                                                                                                                             Carter
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RESULT 14
ADA54961
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC encoding a Propionibacterium acnes protein. The invention also relates to comply peptides encoded by the polynucleotides (ABM35624-ABM6436) and to comply peptides. The invention also relates to compare the comparison of the invention comparison proteins of polypeptides. The invention comprising a golypeptides of the invention, fusion proteins comprising a polypeptide of the invention; an immune response specific for a P. acnes comparison a polypeptide of the invention; a compression proteins comparising a polypeptide of the invention; a compression proteins of polypeptides of the invention; a compression proteins of polypeptides of the invention; a compression proteins of polypeptides of the invention; a compression proteins of polypeptides, compression proteins, and the composition comprising P. acnes polypeptides, compression proteins, T cell specific for a P. acnes in a compression proteins, T cell populations, or compression proteins, T cell populations, or compression proteins, T cell populations or antigen-presenting cells that express the polypeptides, antibodies, fusion compression, and a method for inhibiting the development of P. acnes in a comparison proteins, T cell populations or antigen-presenting cells that express the comparison. The populations or antigen-presenting cells that express the compression proteins, T cell populations or antigen-presenting cells that express the comparison proteins. The polyment of P. acnes in a compression proteins, T cell populations or antigen-presenting cells that express the comparison proteins. The polyment of P. acnes in a comparison proteins, T cell populations or antigen-presenting cells that express the comparison proteins. The polyment of P. acnes in a comparison protein proteins of P. ac
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Best Local Similarity
                                                                                                                                                                                                                                                                                             Cytostatic; Anti-inflammatory; Osteopathic; Neuroprotective; Nootropic; Gene Therapy; human; secretory protein; membrane proteins; cancer; inflammatory disease; osteoporosis; neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                EP1293569-A2
                                                                                                                                                                                                                                                                                                                                                                                    Human protein,
                                                                                                                                                                                                                                                                                                                                                                                                                               20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA54961 standard; protein; 113 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 53 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated polynucleotide (ACF64435-ACF64733) encoding a Propionibacterium acnes protein. The invention also relates
(REAS-)
                                                              14-SEP-2001;
24-JAN-2002;
                                                                                                                             21-MAR-2002;
                                                                                                                                                                        19-MAR-2003.
                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                              2001JP-00328381.
2002US-0350435P.
                                                                                                                               2002EP-00006586.
                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID 2529.
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Pred. No.
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BIOTECHNOLOGY

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RESULT 15
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                                                                                                                                                                                                                                                 31-MAR-1999;
02-APR-1999;
05-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antianaemic; gene therapy; cancer; proliferative disorder; hypertension neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID, AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound; bone damage; cartilage damage; antiinflammatory disease; coagulation; thrombosis; contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; hypotensive; dermatological; antifungal; antirheumatic; antithyroid; ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to novel human secretory or membrane proteins (ADA54072-ADA55710) and their coding sequences (ADA52433-ADA54071). The coding sequences are useful in the gene therapy of diseases caused by abnormalities of the proteins, e.g. cancer, inflammatory diseases, osteoporosis or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polynucleotides encoding full-length polypeptides, e.g. secretory and/or membrane proteins, useful for developing medicines for disease which the gene is involved, or as target molecules for gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human ORFX ORF1791 polypeptide sequence SEQ ID NO:3582.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-2001
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N-PSDB; ADA53322.
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                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-2000; 2000WO-US008621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-OCT-2000.
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                                                                                                                            (CURA-) CURAGEN CORP
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5; Conserv
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                                                                                                                                                                                                              2000US-00540763
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99US-0127636P.
99US-0127728P.
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Hio Y, Otsuka K, Nagai
Otsuka M, Nagahari K, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>.</u>
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Pred. No. 1.1e+02;
0; Mismatches 2;
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Masuho
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K, Irie R,
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DR N-PSDB; AAC76236.

XX

Novel nucleic acids and peptides derived from open reading frame X, proliferative disorders, proliferative, penalogue, proliferative; prolif
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Query Match
Best Local Similarity 71.3%; Score 34; DB 3; Length 138;
Best Local Similarity 71.4%; Pred. No. 1.3e+02;
Matches 5; Conservative 0; Mismatches 2; Indels 0; Gaps

Qy 4 CXPXTGC 10
Db 15 CGPRTGC 21

Bearch completed: January 12, 2005, 23:10:52

Job time: 150.9 secs

0

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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OM protein - protein search, using sw model

Run on: January 12, 2005, 23:04:08 ; Search time 16.8 Seconds (without alignments) 40.090 Million cell updates/sec

Title: Perfect score: Sequence: US-09-932-322-8 38

Scoring table: 1 CXPXTGC 7

BLOSUM62 Gapop 10.0 , Gapext 0.5

283416 seqs, 96216763 residues

Searched:

Total number of hits satisfying chosen parameters: 283416

Minimum DB seg length: 0
Maximum DB seg length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0		Result No. So
0000000	, 	Score
	78.994 78.995 78.995 78.995 78.995 78.995 78.995 78.995 78.995 78.995 78.995	Query Match
1722 1722 119 119 119	1145 1051 1051 1051 1051 1051 1051 1132 1132 1132 1132 1132 1132 1334 1334	
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T23864 E89753 H712662 B45537 S24292 S24294	A27020 R27983 R507983 R504948 F96006 F96006 F96006 F96006 F96006 F96006 F96006 F04331 R24255 AB0331 R24255 AB0331 R24255 AB0331 R149486 R04926 R07729 F04098 F04	ID
12. 13 mil ii	DIF-induced presta phospholipase A2 h hypothetical prote hypothetical prote hypothetical prote s-receptor kinase-hypothetical prote s-receptor kinase-hypothetical prote chorion class A proceonion class A proteonion class A proteonion-induced	ription

ALIGNMENTS

pundas protein precursor - slime mold (Dictyostelium discoideum) (filum discoideum) (filu	Qy 1 CXPXTGC 7 Db 105 CDPETGC 111	Score 34; DB 2; Length 145; Pred. No. 18; 0; Mismatches 2; Indels 0; Gaps	common tiger snake (common tiger snake) (common tiger snake, mainland tiger snak or o7-Sep-1990 #text_change 09-Jul-2004 nnuary 1989 ABL:X14043; NID:g64109; PIDN:CAA32201.1; predicted <sig> atus predicted <mat></mat></sig>	Qy 1 CXPXTGC 7 Db 245 CSPSTGC 251	Score 36; DB 2; Length 601; Pred. No. 23; 0; Mismatches 2; Indels 0; Gaps	- slime mold r-1989 #text_c s.; Mahbubani cestalk gene e MID:3568124 protein #stat
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hypothetical protein F8B4.80 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004 C;Accession: T05348 R;Bevan, M.; Terryn, N.; Ardiles, W.; Buysshaert, C.; Dasseville, R.; De Clerck, ewes, H.W.; Mayer, K.F.X.; Schueller, C. submitted to the Protein Sequence Database, February 1999 A;Reference number: Z15409 A;Reference number: Z15409
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A, Note:
             RiPinan, T.M.; Weidner, S.; Wong, K.; Buhrmester, J.; Chain, P.; Proc. Natl. Acad. Sci. U.S.A. 98, 9889-9894, 2001
A;Title: The complete sequence of the 1,683-kb pSymB megaplasmid A;Reference number: A95842; MUID:21396508; PMID:11481431
A;Accession: F96006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-1058 <BEV>
A;Cross-references: UNIPROT:Q9LEZ5; EMBL:AL357612
A;Experimental source: cultivar Columbia; BAC clone T22D6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Accession: T50496
R;Bevan, M.; Terryn, N.; Ardiles, W.; Buysshaert, C.; Dasseville, ewes, H.W.; Rudd, S.; Lemcke, K.; Mayer, K.F.X. submitted to the Protein Sequence Database, May 2000
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A,Residues: 1-312 <BEV>
A,Cross-references: UNIPROT:Q98UV3; EMBL:AL034567
A,Experimental source: cultivar Columbia; BAC clone
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A, Accession: T50496
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 21-Jul-2000 #sequence_revision 21-Jul-2000 #text_change 09-Jul-2004
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A, Introns: 1/3;
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A;Introns: 60/3; ]
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A;Status: preliminary
                                                                                                                                                                                        hypothetical
                                                                                                                                                                                                                                      RESULT 5
                                                                                                                     ppothetical protein [imported] - Sinorhizobium meliloti (strain 1021) magaplasmid
;Species: Sinorhizobium meliloti
;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 09-Jul-2004
;Accession: P96006
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Best Local
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Best Local
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5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                               86.8%;
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Pred. No. 32;
0; Mismatches
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pred. No. 1.3e+02;
0; Mismatches 2
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A, Cross references: UNIPROT: (092U25; GB:AL591985; PIDN: CAC49718.1; PID: 9
A, Experimental source: strain 1021, megaplasmid pSymB
R;Galibert, F.; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F
pela, D.; Chain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.
L.; Hyman, R.W.; Jones, T.
Science 293, 668-672, 2001
A, Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; K
hebault, P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.;
A;Title: The composite genome of the legume symbiont Sinorhizobium meli
A, Reference number: A99039; MUID: 21368234; pMID: 11474104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        S-receptor kinase-like protein - Arabidopsis thaliana
N;Alternate names: protein F17C15.120
C;Species: Arabidopsis thaliana (mouse-ear cress)
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-643 <BRA>
A;Cross-references: UNIPROT:Q22886; EMBL:U64833; PIDN:AAB04822.1; GSPDB:GN00023; CESP:B
A;Cross-references: UNIPROT:Q22886; EMBL:U64833; PIDN:AAB04822.1; GSPDB:GN00023; CESP:B
                                                                                                                                                                                                                      hypothetical protein B0507.1 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #teC;Accession: T25473 R;Bradshaw, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R; Bevan, M.; Pohl, T.; Weizenegger, submitted to the Protein Sequence Da
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A;Genome: plasmid
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A; Residues: 1-64 < KUR>
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A; Accession: T48397
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A; Note: F17C15.120
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A; Residues: 1-482 <BEV>
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                                                                                                                  A; Status: preliminary; translated
                                                                                                                                        A; Accession:
                                                                                                                                                            A; Description: The sequence A; Reference number: Z20039
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A;Experimental source: cultivar Columbia; BAC clo
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Pred. No. 1
                                                                                                                      from
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Rispoerel, N.; Nguyen, H.T.; Kafatos, F.C.
J. Mol. Biol. 190, 23-35, 1986
A;Title: Gene regulation and evolution in the chorion locus
A;Reference number: A92929; MUID:87060979; PMID:3023635
A;Accession: A24255
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J. Mol. Biol. 190, 23-35, 1986
A/Title: Gene regulation and evolution in the chorion locus
A,reference number: A92929; MUID:87060979; PMID:3023635
A,Recession: B24255
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C,Species: Bombyx mori (silkworm)
C,Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 09-Jul-2004
C,Accession: A24255
sigma E factor regulatory protein rseC [imported] - Yersinia pestis (strain C,Species: Yersinia pestis (strain C,Species: Yersinia pestis C,Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004 C,Accession: AB0331 R,Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; P1
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A; Residues: 1-129 <8PO>
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A;Introns: 59/3; 133/1; 464/3; 586/1
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A; Residues: 1-132 < SPO>
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jl-21/Domain: signal sequence #status predicted <SIG>
jl-21/Domain: signal sequence #status predicted <SIG>
j22-132/Product: chorion class A protein L12 #status predicted <MAT>
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Pred. No. 89;
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Pred. No. 90;
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C;Genetics:
A;Gene: rseC
C;Superfamily: Escherichia coli sigma-E factor regulatory protein rseC
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A; Map position: 6
C; Superfamily: Neurospo
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C;Superfamily: hevein precursor; barwin homology; hevein chitin-b F;1-25/Domain: signal sequence #status predicted <SIG> F;26-200/Product: wound-induced protein 1 #status predicted <MAT> F;26-69/Domain: hevein chitin-binding domain homology <HCB>
                                                                                                                                                                                                                                                      R;Stanford, A.; Bevan, M.; Northcote, D. Mol. Gen. Genet. 215, 200-208, 1989
A;Title: Differential expression within a fa;Reference number: S04926; MUID:89218921;
                                                                                                                                                                                                                                                                                                                                              wound-induced protein 1 precursor - potato
C;Species: Solanum tuberosum (potato)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 09-Jul-2004
C;Accession: S04926
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A;Residues: 1-199 <SCH>
A;Croes-references: EMBL;AL356173; GSPDB;GN00116; NCSP:B14D6.380
A;Experimental source: BAC clone B14D6; strain OR74A
C;Genetics:
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A; Accession: T49486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Accession: T49486
R;Schulte, U.; Aign, V.; Hoheisel, J.; Brandt, P.; Fartmann, B.; Holland, R.; submitted to the Protein Sequence Database, May 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein B14D6.380 [imported] - Neurospora crassa
C;Species: Neurospora crassa
C;Date: 02-Jun-2000 #sequence_revision 02-Jun-2000 #text_change 18-Aug-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-151 < KUR>
                                                                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-200 <STA>
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                                                                                            A; Introns: 143/3
                                                                                                                A; Gene: win1
                                                                                                                                           C;Genetics:
                                                                                                                                                              A;Cross-references: UNIPROT:P09761; EMBL:X13497; NID:g21617; PIDN:CAA31851.1;
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4; Conserve
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57.1%;
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1; Mismatches
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Pred. No. 1e+02;
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potato

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RESULT 15
S04927
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C;Bpecies: Lycopersicon esculentum (tomato)
C;Date: 14-May-1999 #sequence_revision 14-May-1999 #text_change 09-Jul-2004
C;Accession: T07729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Harris, N.; Taylor, J.E.; Roberts, J.A.
J. Exp. Bot. 48, 1223-1227, 1997
A;TItle: Characterization and expression of an mRNA encoding a wound-induced (win) prote A;Reference number: 216099
A;Accession: T07729
A;Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                A/Title: A novel pathogen- and wound-inducible tobacco A/Reference number: Z15209; MUID:94159785; PMID:8115541 A/Accession: T04098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CBP20 preproprotein - common tobacco
N;Alternate names: wound-induced protein
C;Species: Nicotiana tabacum (common tobacco)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:Q41231; EMBL:S72452; NID:g632733; PIDN:AAB29959.1; PID:g6327A;Experimental source: cultivar Samsun NN C;Superfamily: hevein precursor; barwin homology; hevein chitin-binding domain homology F;23-66/Domain: hevein chitin-binding domain homology <ACB>
                                                                                                                                                                                                                                                                                                                                                                                               A,8tatus: preliminary; translated from GB/EMBL/DDBJ
A,Molecule type: mRNA
A,Residues: 1-211 <PON>
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A, Residues: 1-201 <HAR>
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  wound-induced
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Plant Physiol. 104, 109-118, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F;68-188/Domain: barwin homology <BAR>
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4; Conservative
protein 2 precursor - potato
                                                                                                             CSPSQGC 59
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Pred. No. 1.2e+02;
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Pred. No. 1.3e+02;
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RiStanford, A.; Bevan, M.; Northcote, D.
Mol. Gen. Genet. 215, 200-208, Mol. Gen. Genet. 215, 200-208, Title: Differential expression within a family of novel wound-induced A;Title: Differential expression within a family of novel wound-induced A;Reference number: S04926; MUID:89218921; PMID:2710099
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Solanum tuberosum (potato)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 09-Jul-2004
C;Accession: S04927
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                                                                                                                                                                                                    C;Superfamily: hevein precursor; barwin homology; hevein chitin-binding domain homology F;1-25/Domain: signal sequence #status predicted <SIG>F;26-211/Product: wound-induced protein 2 #status predicted <MAT>F;26-69/Domain: hevein chitin-binding domain homology <HCB>F;77-198/Domain: barwin homology <BAR>
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A; Residues: 1-211 <STA>
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                                                                                                  Similarity 57.1
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57.1%;
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Search completed: January 12, 2005, 23:16:03 Job time : 16.8 secs

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 12, 2005, 22:56:08; Search time 88.55 Seconds (without alignments) 45.484 Million cell updates/sec

US-09-932-322-8 38

Title: Perfect score: Sequence: 1 CXPXTGC 7

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 1825181 seqs, 575374646 residues

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2 Q91ERB 2 Q228B6 2 Q7N375 2 P90569 2 P90569 2 P771300 1 CHA1 BOMMO 2 Q8MQ78 2 Q89958 1 CHA2 BOMMO 1 CHA2 BOMMO 2 Q9LIW4 2 Q8E016 2 Q8E016 2 Q8E016 2 Q8E016 2 Q8E016	78.9	78.9	78.9	78.9	78.9	78.9	78.9	78.9	78.9	78.9	78.9	78.9	81.6	81.6
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	Aas62	Q8zd76	Q6d216	Q9liw4	P08825	P08826	856480	Q8mq78	P08929	P77130	P90569	Q7n375	Q22886	Q9lzr8

ALIGNMENTS

Query M Best Lo Matches			RESULT 1 Q98AR9 AC Q98 DT 01- DT 01- DT 01- CGN O1- GGN O1- GG
atch cal	SEQUENCE FROM N.A. STRAIR-MAFF30309; MEDLINE=21082930 Kaneko T., Nakamur Watanabe A., Idesa Mishida Y., Kiyoka Mishida Y., Naka Takeuchi C., Yamad "Complete genome s Mesorhizobium loti DNA Res. 7:331-338 EMBL; RAP003007; EM EMBL; RAP003007; EM EMBL; RAP003007; EM EMBL; RAP003017; FR GO; GO:0016491; F: GO; GO:0016491; F: GO; GO:0016491; F: GO; FR00015; P: InterPro; IPR00234 Pfam; PF000061; adh PRINTS; PR000061; Complete proteome. SEQUENCE 192 AA;	SEQUENCE FROM N.A. STRAILMARF303099; MEDLINE=2108295; Kaneko T., Nakamux Watanabe A., Idesa Kishida Y., Kiyoka Mochizuki Y., Naka Mochizuki Y., Naka Mochizuki O., Yamad "Complete genome some some leti DNA Res. 7:381-406	LT 1 R9 R9 R9 R9 R9 R1 O1-OCT-2001 (TrE O1-OCT-2001 (TrE O1-EXECUTE (TrE O1-EXECUTE (TrE O1-EXECUTE (TrE M1r5880 protein. OrderedLocusName Rhizobium loti (Bacteria, proteo Bryllobacteriace NCBI_TAXID=381;
94. Similarity 71. 5; Conservative	SEQUENCE FROM N.A. STRALN=MAPF303099; STRALN=MAPF303099; MEDLINE=21082930; PubMed=11214968; MEDLINE=21082930; PubMed=11214968; MEDLINE=21082930; PubMed=11214968; Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsumoto Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimot. Takeuchi C., Yamada M., Tabata S.; "Complete genome structure of the nitrogen-fixing symbioti. Mesorhizobium loti."; Meso	SEQUENCE FROM N.A. STRALINE—MARF30309; STRALINE—21082936, PubMed=11214974; MEDLINE=21082936, PubMed=11214974; MEDLINE=21082936, PubMed=11214974; MATERIAL TO S., Asamizu E., Kato S., Asamizu E., Kato T., Kawashima K., Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Takeuchi C., Yamada M., Tabata S.; Takeuchi C., Yamada M., Tabata S.; Takeuchi C., Yamada M., Tabata S.; Mesorhizobium loti (supplement)."; DNA Res. 7:381-406(2000).	T 1 99 AP PRELIMINARY; PRT; 192 AA. 998AR9 PRELIMINARY; PRT; 192 AA. 998AR9; 01-OCT-2001 (TYEMBLrel. 18, Created) 01-OCT-2001 (TYEMBLrel. 26, Last sequence upda 01-MAR-2004 (TYEMBLrel. 26, Last annotation up M1r5880 protein. OrderedLocusNames=m1r5880; Rhizobium loti (Mesorhizobium loti). Bacteria; Proteobacteria; Alphaproteobacteria; Phyllobacteriaceae; Mesorhizobium. NCBI_TaxID-381; [1]
4 %	N.A. 3099; PubMed=11214968; 930; PubMed=11214968; 1desawa K., Ishikawa iyokawa C., Kohara M. Nakayama S., Nakazak Yamada M., Tabata S.; Ome structure of the 10ti."; 1-336(2000). 1-336(2000). 7; BAB52253.1; 7; BAB52253.1; 1FK8. 11FK8. 11FK8. 11FK8. 12; Pontidoreductase a 1; F:oxidoreductase a 2; P:mediatase a 2; P:mediatase a 3; F:oxidoreductase a 4; F:oxidoreductase a 4; F:oxidoreductase a 5; F:oxidoreducta	Med=1121, Sato (K., Ishil C., Kohas a S., Nal i, Tabati cuture of cuture of cutupplement	RY; 1. 18, C; 1. 26, L; 1. 26, L; 5880; hizobium ria; Alph lesorhizol
Score 36; DB Pred. No. 27; Nismatches	4968; S., Asamizu S., Asamizu Raw A., Matsu kazaki N., i a S.; the nitroge tt. IEA. tt_C2.	p ⊢. > ø	PRT; 192 ; Created) Last sequence Last annotati Last annotati in loti). lphaproteobact. zobium.
, DB 2; . 27; :ches	68; ABSAMIZU E., KATO T., SABAMO WA A., KAWASHIMA K., Kimura T M., MATSUMOTO M., MATSUMO A. zaki N., Shimpo S., Sugimoto S.; he nitrogen-fixing symbiotic he activity; IEA. EA. C2. FD2F660D156037BC CRC64;	amizu E., Kato T., Sasamot ., Kawashima K., Kimura T. Matsumoto M., Matsuno A., N., Shimpo S., Sugimoto M.	r; 192 AA. sequence update) annotation update) ci). coteobacteria; Rhiz
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Q86AK7;
01-JUN-2003
01-JUN-2003
01-MAR-2004
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Dictyostelium discoideum (Slime mold).
Eukaryota, Mycetzzoa, Dictyosteliida, Dictyostelium
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OBKJ72; PRELIMINARY; PRT; 546 AA.

OBKJ72; OFFICE OF OFFICE OFFICE
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EMBL; AC116994; AA051380.1; -

HSSP; P01382; INTN

InterPro; IPR002172; LDL_receptor_A.

InterPro; IPR001673; S. mold repeat.

Pfam; PF00526; D1cty_CTDC; I3.

ERINTS; PR00361; LDLRECEPTOR.

SEQUENCE 360 AA; 38013 MW; 480061ABA26EDB1E CRC64;
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                                                                                           MEDILINE-21999272; PubMed=12003951;
Sullivan J.T., Trzebiatowski J.R., Cruickshank R.W., Gouzy J.,
Brown S.D., Ellitot R.W., Fleetwood D.J., McCallum N.G., Rossbach U.,
Stuart G.S., Weaver J.B., Webby R.J., de Bruijn F.J., Ronson C.W.;
"Comparative sequence analysis of the symbiosis island of
Mesorhizobium loti strain R7A."
J. Bacteriol. 184:3086-3095(2002).
                                                                                                                                                                                                                                                                                                                                                                               Rhizobium loti (Mesorhizobium loti).
Bacteria, Proteobacteria, Alphaproteobacteria,
Phyllobacteriaceae, Mesorhizobium.
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                                             SIMILARITY: Belongs to (SDR) family.
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77114; CAD31361.1;
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(TrEMBLrel. 24, Last sequence update)
(TrEMBLrel. 26, Last annotation update)
Dictyostelium discoideum (Slime mold). P
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71.4%;
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Q7M4J3;
Q1-MAR-2004
01-MAR-2004
01-MAR-2004
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InterPro; IPR002198; ApH_short.
InterPro; IPR002347; Adh_short_C2.
Pfam; PF00106; adh short; 2.
PRINTS; PR00080; GDHRDH.
PRINTS; PR00080; SDRFAMILY.
PROSTE; P800080; SDRFAMILY.
PROSTE; P8000861; ADH_SHORT; 1.
                                                                                      01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protein.
Macaca fascicularis (Crab eating macaque) (Cynomolgu Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Cercopithe Cercopithecinae; Macaca.
                                                                                                                                                                                              Q9N018;
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01-MAR-2004 (TrEMBLrel. 26, Last expense update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DIF-induced prestalk pDd63 protein precursor (Fragments).
Dictyostelium discoideum (Slime mold).
Bukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.
 SEQUENCE FROM N.A.
TISSUB-Cerebellum cortex;
TISSUB-Cerebellum cortex;
Osada N., Hida M., Kusuda J., Tanuma R., Iseki K., Hirai
Osada N., Sugano S., Hashimoto K.;
Suzuki Y., Sugano S., Hashimoto K.;
Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                             Pfam; P
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PIR; A27020; A27020.
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Early A., Berks M., Jermyn K.A.;
"Direct induction of Dictyostellum prestalk gene expression
provides evidence that DIF is a morphogen.";
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Pfam; PF00526; Dicty_CTDC; 19.
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71.48;
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ertebrata; Euteleostomi;
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Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Butheria; Primates; Catarrhini; Cercopithecidae;

Cercopithecinae; Macaca.
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Q95KD2; PRELIMINARY;
Q1-DEC-2001 (TrEMBLrel. 19,
Q1-DEC-2003 (TrEMBLrel. 25,
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Hypothetical protein.
SEQUENCE 148 AA; 15495 MW
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QPAQPS;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Putative ABC transporter subunit.

Name=ORF31;
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TISSUE-Medulla oblogata;
TOSUE-Medulla oblogata;
THISSUE-MEDIA
THICA M., Kusuda J., Tanuma R., Iseki
Suzuki Y., Sugano S., Hashimoto K.;
Submitted (JUN-2001) to the EMBL/GenBank/DDBJ
EMBL, AB062948; BAB60737.1; -.
                                                                                                                                                       MEDLINE=97386424; PubMed=9244273; Sato S., Ouchiyama N., Kimura T., Nojiri H., Yamane H., Omori T.; Sato S., Ouchiyama N., Kimura T., Nojiri H., Yamane H., Omori T.; "Cloning of genes involved in carbazole degradation of Pseudomonas strain CAlO: nucleotide sequence of genes and characterization of meta-cleavage enzymes and hydrolase."; J. Bacteriol. 179:4841-4849(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                           Pseudomonas resinovorans.
Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
  MEDLINE=97386425; PubMed=9244274;
Sato S., Nam J., Kasuga K., Nojiri H.,
"Identification and characterization o
                                                                                                                                                                                                                                                                                                                       STRAIN-CA10
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S., Hashimoto K.;
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Pred. No. 33;
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  Yamane H., Omori T.; f genes encoding carbazole 1,
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EMBL; AB047548; BAB32742.1; -.

GO; GO:0016020; C:membrane; IEA.

GO; GO:0005215; F:transporter activity;

GO; GO:0005810; P:transport; IEA.

InterPro; IPR001851; Bac_inmem_transp.

InterPro; IPR001851; Ribosomal S2.
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Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; BC053646; AAH53646.1; -.
Hypothetical protein.
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Mammalia; Eutheria; Primates;
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"Genetic characterization and evolutionary implications of car gene cluster in carbazole-degrader, Pseudomonas sp. strain CA10.";
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TISSUE=Lung;
MEDLINE=22388257; PubMed=12477932;
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J. Bacteriol. 179:4850-4858(1997).
                                                                                                                                                                                                                                                                                                                                                                        and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A.
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           PROSITE;
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ACT_SITE
DISULPID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-FEB-1991 (Rel. 17, Last sequence update)
01-OCT-2004 (Rel. 45, Last annotation update)
Probable phospholipase A2 precursor (EC 3.1.1.4) (Phosphatidylcholine
                                                                                                                                                                                                                       Pfam, PF00068; Phospholip A2 1; 1.
PRINTS; PR00389; PHPHLIPASEA2.
Probom; PD000303; PhospholipaseA2;
SMART; SM00085; PA2c; 1.
                                                                                                                                                                                                                                                                           EMBL; X14043; CAA32201.1; -.
PIR; 807983; 807983.
HSSP; P00608; 1AE7.
InterPro; IPR001211; PhospholipaseA2.
                                                                                                                                                                                                                                                                                                                                             use by non-profit institutions as long as modified and this statement is not removed. It entities requires a license agreement (See htt or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no wa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Notechia scutatus scutatus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blapidae, Acanthophiinae,
                                                                                                                                                                                              PROSITE; PS00119; PA2 ASP; 1.
PROSITE; PS00118; PA2 HIS; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mitted (JAN-1989) to the EMBL/GenBank/DDBJ databases. FUNCTION: PAZ catalyses the calcium-dependent hydrolysis acyl groups in 3-sn-phosphoglycerrides. CATALYTIC ACTIVITY: Phosphatidylcholine + H(2)O = 1-caryllycerophosphocholine + a carboxylate. CATALYTIC Binds I calcium ion per subunit (By similarity) SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      subfamily.
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5; Conserv
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e; Notechis.
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0; Mismatches
         Probable phospholipase A Probable phospholipase A By similarity. By similarity.
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              (via carbonyl oxygen)
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                      Q9SUV3;
Q9SUV3;
01-MAY-2000
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Q7U7V9; O1-OCT-2003 (TrEMBLrel. 25,
01-OCT-2003 (TrEMBLrel. 25,
01-OCT-2003 (TrEMBLrel. 25,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                METAL
                                                                                  Bevan M., Terryn N., Ardiles W., Buyss
De Clerck R., De Keyser A., Neyt P., R.
Villaroel R., Gielen J., Van Montagu M
Mayer K.F.X., Lemcke K., Schueller C.,
Submitted (DEC-1998) to the EMBL/GenBa
                                                                                                                                                                                                                                                                      Hypothetical protein F8B4.80 (Hypothetical protein AT4g32380).

Name=F8B4.80; Synonyms-AT4g32380;

Arabidopsis thaliana (Mouse-car cress).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheoph
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; r
spermatophyta; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

MEDLINE=22825697; PubMed=12917641; DOI=10.1038/nature01943;
Palenik B., Brahamsha B., Larimer F.W., Land M.L., Hauser L.
Chain P., Lamerdin J.E., Regala W., Allen E.E., McCarren J.
Paulsen I.T., Dufresne A., Partensky F., Webb E.A., Waterbu
"The genome of a motile marine Synechococcus.";
Nature 424:1037-1042(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synechococcus sp. (strain WH8102).
Bacteria; Cyanobacteria; Chroococcales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OrderedLocusNames=SYNW0871;
SEQUENCE FROM N.A.

EU Arabidopsis sequencing project;
Submitted (AUG-1999) to the EMBL/GenBank/DDBJ
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                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                   NCBI_TaxID=3702;
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; 38E36029AEE5FAC9 CRC64;
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                                                                                                                                Buysshaert C., Da
P., Rouze P., Var
tagu M., Hoheisel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               312
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Van
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    databases
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                                                                                                                                                                Dasseville R.,
Van Den Daele H.,
                                                                                                                                        J.
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RESULT 12
Q6TVH9
ID V6TVH
AC Q6TVH
DT 05-JU
DT 0
RESULT 13
AAR98366
ID AAR98
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GO; GO:0005975; P:carbohydrate metabolism; IEA.
InterPro; IPR000743; Glyco_hydro_28.
InterPro; IPR006626; PbH1.
InterPro; IPR011050; Pecttin_lyas_like.
Pfam; PP00295; Glyco_hydro_28; 2.
SMART; SM00710; PbH1; 3.
Cell_wall; Glycosidase; Hydrolase; Hymothotical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Genomes of the Parapoxviruses Stomtatitis Virus.";
J. Virol. 78:168-177(2004).
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InterPro; IPR007027; Pox F11.
Pfam; PF04943; Pox F11; 1.
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   AAR98366
                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical
SEQUENCE 4
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de la Concha-Bermejillo
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465 AA; 5
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312 AA; 3409
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   PRELIMINARY;
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71.4%;
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      PRT;
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2;
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CRC64;
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RESULT 14
Q8LJ36
SOUTH THE PROPERTY OF THE PROP
       RA Sasaki T., Matsumoto T., Yamamoto K., Sakata K., Baba T., Katayose Y., RA Wu J., Niimura Y., Cheng Z., Nagamura Y., Antonio B.A., Kanamori H., RA Hosokawa S., Masukawa M., Arkawa K., Chiden Y., Hayashi M., Okamoto M., Ando T., Aoki H., Arita K., Hamada M., Harada C., RA Hijishita S., Honda M., Ichikawa Y., Idonuma A., Iijima M., Ikeda M., RA Ikeno M., Itoh S., Itoh T., Itoh Y., Itoh Y., Iwabuchi A., Kamiya K., RA Karasawa W., Katagiri S., Kikuta A., Kobayashi N., Kono I., RA Machita K., Maehara T., Mizuno H., Mizubayashi N., Kono I., Nagasaki H., Nakashima M., Nakama Y., Nakamichi Y., Nakamura M., Nagasaki H., Nakashima M., Nakama Y., Nakamichi Y., Nakamura M., Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K., RA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K., RA Zhong H., Iwama H., Zmdo T., Ito H., Hahn J.H., Kim H.I., Eun M.Y., RA Yano M., Jiang J., Gojobori T.; The genome sequence and structure of rice chromosome 1."; Nature 420.312-316(2002).

DR EMBL; AP003344; BAC07361.1; -. DR Go; Go:0004194; F:pepsin A activity; IEA. Go; Go:0004194; F:pepsin A activity; IEA. DR Go; Go:0006508; P:proteolysis and peptidolysis; IEA. DR Go; Go:0006508; Peptidase Al, Entrys; Prantys; Prantys; Peptidase Al, Peptims; Pro0026; Asp; 1.
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Matches 5
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Q8LJ36;
01-OCT-2002
01-OCT-2002
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02-MAR-2004
02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oryza sativa (japonica cultivar-group).
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Enrhartoideae; Oryzeae; Oryza.
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01-OCT-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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J. Virol. 78:168-177(2004).
EMBL; AY386265; AAR98366.1; -.
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Bovine papular stomatitis virus.
Viruses; dsDNA viruses, no RNA s
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Name=P0413G02.18;
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de la Concha-Bermejillo
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465 AA; 5
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D., Piccone M.E.,
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RESULT 15
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Search completed: January 12, 2005, 23:15:11 Job time : 88.55 sece
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OB69K4,

O1-JUN-2003 (TrEMBLrel. 24, Created)

O1-JUN-2003 (TrEMBLrel. 24, Last sequence update)

O1-MR-2004 (TrEMBLrel. 26, Last annotation update)

Similar to Mus musculus (Mouse). Tenascin X.

Dictyostelium discoideum (Slime mold).

Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Baumgart C.;

Baumgart C.;

Bubmitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; AC115598; AAO53206.1; -.

HSSP; P01132; 1EGF.

GO; GO:0005198; F:structural molecule activity; IEA.

InterPro; IPR000742; EGF 1;

InterPro; IPR006210; IEGF 1;

InterPro; IPR006210; IEGF.

InterPro; IPR006210; IEGF.
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PRINTS; PR00011; EGF; 9.

SMART; SM00181; EGF; 9.

PROSITE; PS01022; EGF 1; UNKNOWN_10.

PROSITE; PS01186; EGF 2; 7.

PROSITE; PS50026; EGF 3; 6.

PROSITE; PS50026; EGF 3; 6.
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MEDLINE-22092622; PubMed=12097910;
Gloeckner G., Eichinger L., Szafranski K., Pachebat J., Dear P.,
Gloeckner G., Baumgart C., Parra G., April J.F., Guigo R., Kumpf K.,
Tunggal B., Cox E., Quail M.A., Platzer M., Rosenthal A., Noegel A.A.;
"Sequence and analysis of chromosome 2 of Dictyostelium discoideum.";
Nature 418:79-85(2002).
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STRAIN=AX4;
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                                                                                                                                                                                                                                                                                            Conservative
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Perfect score:
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Match
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38
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Listing first 45 summaries
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138
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AAY04992	AAY04996	AAY39105	AAY38968	AAW81666	AAW64303	AAW32431	AAW32363	ABB93340	ABO70819	ABO74638	ABP76713	ABP99348	ABP99259	ADF60298	AAC00015	ADF58925	ADJ71040	ABP06368	AB081251
Aay04992		Aay39105	Aay38968	Aaw81666	Aaw64303	Aaw32431	Aaw32363	Abb93340	Abo70819	Abo74638	Abp76713	Abp99348 (Abp99259 (Adf60298 1	Aao00015 1	Adf58925 1	Adj71040	Abp06368 1	Abo81251
Mycobacte	Mycobacte	M. tuberc	M. tuberc	M. tuberc	Mycobacte	Mycobacte	Mycobacte	Herbicida	Pseudomon	Pseudomon	Streptomy	Orthosomy	Orthosomy	Human con	Human pol	Human pol	Human hea	Human ORF	Pseudomon

ALIGNMENTS

SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant. AAU49413; AAU49413 standard; protein; 79 WO200181581-A2 Propionibacterium acnes immunogenic protein #10309 13-FEB-2002 Propionibacterium acnes. (first entry) Ā

×	MOZOGIOISGI-AZ:	
PD	01-NOV-2001.	
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PF	20-APR-2001; 2001WO-US012865.	
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PR	21-APR-2000; 2000US-0199047P.	
PR	02-JUN-2000; 2000US-0208841P.	
PR	07-JUL-2000; 2000US-0216747P.	
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PA	(CORI-) CORIXA CORP.	
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Id	Skeiky YAW, Persing DH, Mitcham JL, Wang SS, Bhatia A;	
Ιď	L'maisonneuve J, Zhang Y, Jen S, Carter D;	
ğ		
DR	WPI; 2001-616774/71.	
DR	N-PSDB: AAS59545.	

Example 1; SEQ ID NO 10608; 1069pp; English.

Propionibacterium acnes vaccinating against and treating acne vulgaris.

polypeptides and nucleic acids useful for diagnosing infections, especially useful

for

Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory

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ABM4592

ABM4592

ABM4592

ABM4

ACC ABM4

ACC
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Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitcham Chang Y,
encoding a Propionibacterium across protein. The invention also relates to polypeptides encoded by the polynuclectides (ABM/35624 ABM/64536) and to immunogenic fragments of P. across polypeptides. The invention additionally encompasses expression vectors and host cells comprising a polynuclectide of the invention; antibodies against polypeptides of the invention; a method for stimulating an immune response specific for a P. across polypeptide and an isolated T cell population comprising T cells prepared via this method; a vaccine composition (comprising P. across polypeptides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
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                                                                                                                                                                                                                                                                                                                                                                                             Example 1, SEQ ID NO 10608; 1481pp; English.
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                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003-381789/36.
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Benson DR,
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Jones R, Carte
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC polynucleotides, antibodies, fusion proteins, T cell populations, or cantigen-presenting cells that express the polypeptide); a method and kit cf for detecting or determining the presence or absence of P. acnes in a cc patient; and a method for inhibiting the development of P. acnes in a cc patient. The P. acnes polypeptides, polynucleotides, antibodies, fusion proteins, T cell populations or antigen-presenting cells that express the cc polypeptides are useful for diagnosing, preventing or treating acnes cc vulgaris, or for stimulating an immune response specific for a P. acnes cc vulgaris, or for stimulating an immune response specific for the acnes cc protein. The polynucleotides can also be used as probes or primers for nucleic acid hybridisation. The vaccine composition is useful for the stimulation of an immune response against P. acnes, or for treating acne, can the kit is useful for performing a diagnostic assay. The present cc and the kit is useful for performing a diagnostic assay. The present cc reading frame) contained within the P. acnes polynucleotides of the cinvention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly cc from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 5
                                                                                                                                                                              Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of
                                                                                                                                                                                                                            N-PSDB;
                                                                                                                                                                                                                                                                        Rubenfield MJ,
                                                                                                                                                                                                                                                                                                                                 18-FEB-1998;
27-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                             18-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                      US6551795-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pseudomonas aeruginosa polypeptide #6603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AB074428;
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                                                                                                                                                                                                                                                                                                     (GENO-) GENOME
                                                                                                                                                                                                                                         2003-615309/58
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  infection; Pseudomonas aeruginosa infection; antibacterial.
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                                                                                                                                                                                                                                                                       Nolling J,
                                                                                                                                                                                                                                                                                                                                 98US-0074788P
98US-0094190P
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71.4%;
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The invention relates to Resudomonas aeruginosa polypeptides and the polymuclectides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a

Disclosure; SEQ ID NO 23174; 455pp; English. pathological conditions resulting from bacterial

bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant

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                                                                                                                                                                                                                                                                                    05-MAY-1998;
06-NOV-1998;
20-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modulation; nonclassical cadherin mediated cell adhesion; CAR; inhibition; cadherin extracellular domain; cell adhesion recognition; OB-cadherin-1; cadherin-5; cadherin-6; cadherin-7; cadherin-8; cadherin-12; cadherin-14; cadherin-15; T-cadherin; PB-cadherin; cadherin related neuronal receptor; LI-cadherin; protocadherin; desmoglein; desmocollin; calcium binding; cancer; tumour; obesity; rheumatoid arthritis; multiple sclerosis; diabetes; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            production of P. aeruginosa-derived peptides or polypeptides, as target components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences of Pseudomonas species using biochip technology. Sequences ABO67826-ABO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at
                                                                                                                    New cadherin modulating agents, used for modulating nonclassical cadherin -mediated functions for treating e.g. cancers, obesity, rheumatoid arthritis, multiple sclerosis, diabetes or a neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAR-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequata.uspto.gov/sequence.html
The present invention describes cadherin modulating agents comprising peptides which comprise a nonclassical cadherin recognition (CAR) sequence. The MAs can be used for modulat nonclassical cadherin-mediated functions. They can be used inhibiting adhesion of nonclassical-cadherin expressing cel
                                                                                                                                                                                                                                                                                                                                               05-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                             11-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PB-cadherin
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                                                                                                                                                                                                                                         (ADHE-) ADHEREX TECHNOLOGIES INC
                                                                                                                                                                               2000-038791/03.
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5; Conserv
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                                                                                                                  functions;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell adhesion recognition cyclic peptide SEQ ID NO:4047
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                                                                                                                                                                                                           Gour BJ,
                                                                                                                                                                                                                                                                     98US-00073040.
98US-00187859.
99US-00234395.
99US-00264516.
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71.4%;
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                                                                                      English
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Pred. No.
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2;
   They can be used for 
rin expressing cells :
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                               modulating
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RESULT 5
AAXYC 1489
AAY C1489
AAY C1489
AAY CAAYG
AX AAYG
AX AAYG
AX CAAH
AX C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
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06-NOV-1998;
20-JAN-1999;
08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OB-cadherin; cadherin-5; cadherin-6; cadherin-7; cadherin-8; cadherin-14; cadherin-15; T-cadherin; PB-cadherin; protocadherin; cadherin related neuronal receptor; LI-cadherin; protocadherin; desmocollin; calcium binding; cancer; tumour; obesity; rheumatoid arthritis; multiple sclerosis; diabetes; metastasis; neurological disease; cyclic
                                                                                                                                                                                                                                                                                                                                                                                                 05-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modulation; inhibition;
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                                                                                                                         Blaschuk OW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9957149-A2
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                                                                                                                                                                                     (ADHE-) ADHEREX TECHNOLOGIES INC
                                                               2000-038791/03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nonclassical cadherin mediated cell adhesion; CAR; cadherin extracellular domain; cell adhesion recognition; cadherin-8; cadherin-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                         Gour BJ,
                                                                                                                                                                                                                                           98US-00073040.
98US-00187859.
99US-00234395.
99US-00264516.
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1. .7
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Pred. No. 1.7e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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New cadherin

modulating

agents, used for modulating nonclassical cadherin

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RESULT 6
AAY62007
ID AAY66
AX AAY66
AX AAY66
AX AAY66
AX AAY66
AX CACH
DT 02-M
AX CACH
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modulation; nonclassical cadherin mediated cell adhesion; CAR; inhibition; cadherin extracellular domain; cell adhesion recognition; OB-cadherin; cadherin-15; cadherin-6; cadherin-7; cadherin-8; cadherin cadherin-14; cadherin-15; T-cadherin; PB-cadherin; cadherin; cadherin related neuronal receptor; II-cadherin; protocadherin; desmocylein; desmocylein; calcium binding; cancer; tumour; obesity; rheumatoid arthritis; multiple sclerosis; diabetes; metastasis; neurological disease; cyclic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cadherin-12 cell adhesion recognition cyclic peptide SEQ ID NO:1799.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -mediated functions arthritis, multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY62007 standard; peptide; 7
    05-MAY-1998;
                                                                                                                              05-MAY-1999;
                                                                                                                                                                                                                 11-NOV-1999.
                                                                                                                                                                                                                                                                                           WO9957149-A2
                                                                                                                                                                                                                                                                                                                                                                                  Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CXPXTGC 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ξ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
        98US-00073040.
98US-00187859.
                                                                                                                              99WO-CA000363
                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
1. .7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for treating e.g. cancers, obesity, rheumatoid sclerosis, diabetes or a neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89.5%;
71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention
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Pred. No. 1.7e+06;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ν
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cell adhesion
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New cadherin modulating agents, used for modulating nonclassical cadherin -mediated functions for treating e.g. cancers, obesity, rheumatoid arthritis, multiple sclerosis, diabetes or a neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JAN-1999;
08-MAR-1999;
                                                                                                                                                                                                                                              cadherin related neuronal receptor; LI-cadherin; protocadherin; desmoglein; desmocollin; calcium binding; cancer; tumour; obesi rheumatoid arthritis; multiple sclerosis; diabetes; metastasis;
                                                                                                                                                                                                                                                                                                                                                                        Modulation; nonclassical cadherin mediated cell adhesion; CAR; inhibition; cadherin extracellular domain; cell adhesion recognition; OB-cadherin-1; cadherin-5; cadherin-6; cadherin-7; cadherin-8; cadherin-16; cadherin-14; cadherin-15; T-cadherin; PB-cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY62224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         recognition (CAR) sequence. The MAS can be nonclassical cadherin-mediated functions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising peptides which comprise a nonclassical cadherin cel recognition (CAR) sequence. The MAs can be used for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 48; Page 180; 252pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-038791/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cadherin-14 cell adhesion recognition cyclic peptide SEQ ID NO:3971.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY62224 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes cadherin modulating
                                                                                                                                                                                                           neurological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDPKTGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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99US-00264516
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 34; DB 3;
Pred. No. 1.7e+06;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              They can be used for e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ng agents (MA)
cadherin cell
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                                                                                                                                                                                                                                                                                                       obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                            cadherin-12;
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sapiens

Location/Qualifiers

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RESULT 8
ABJO ID ABJO
XX ABJO
XX ABJO
XX DT 05-S
XX DE B ly
XX B ly
XX B lift
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC angiogenesis in a mammal, inducing apoptosis in a nonclassical cadherin-
CC expressing cell, preventing or treating obesity in a mammal, stimulating
CC central nervous system, treating a demyelinating drug delivery to the
CC central nervous system, treating a demyelinating neurological disease,
CC increasing vasopermeability in a mammal, enhancing adhesion of
CC nonclassical cadherin-expressing cells, inhibiting synaptic stability in
CC a mammal, or preventing pregnancy in a mammal. They can also be used for
CC e.g. enhancing or directing neurite outgrowth, facilitating wound healing
CC or reducing scar tissue, or enhancing adhesion of foreign tissue in a
CC mammal. They can also be used for treating e.g. psoriasis, arthritis, age
CC erelated macular degeneration, multiple sclerosis and diabetes. The
CC products can also be used for detection and diagnosis and in bioreactors.
CC AAY60592 to AAY6492 represent specifically claimed apptides, and
CC AAY64573 to AAY64643 and AAZ33183 to AAZ33186 represent sequences used in
CC the exemplification of the present invention
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Best Local S
Matches 5
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06-NOV-1998;
20-JAN-1999;
08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New cadherin modulating agents, used for modulating nonclassical cadherin -mediated functions for treating e.g. cancers, obesity, rheumatoid arthritis, multiple sclerosis, diabetes or a neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising peptides which comprise a nonclassical cadherin cell adhesion recognition (CAR) sequence. The was can be used for modulating nonclassical cadherin-mediated functions. They can be used for e.g. inhibiting adhesion of nonclassical-cadherin expressing cells in a mammal, enhancing delivery of a drug through the skin of a mammal, enhancing delivery of a drug to a tumour in a mammal, inhibiting metastasis of a cancer in a mammal, inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9957149-A2
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B lymphocyte stimulator protein binding protein; BLyS; immune disease; allergy; proliferative disease; infectious disease; arteriosclerosis; inflammatory disorder; hypergammaglobulinaemia; blood clotting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 54; Page 184; 252pp; English.
                                                                                                                      05-SEP-2002
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                                                                                                                                                                                                ABJ00550
                                                                             lymphocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention describes cadherin modulating
                                                                                                                                                                                                                                                                                                                                                                     Similarity
5, Conser
                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                 CXPXTGC 7
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                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                           stimulator protein binding peptide #1.
                                                                                                                      (first entry)
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98US-00187859.
99US-00234395.
99US-00264516.
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                                                                                                                                                                                              peptide;
                                                                                                                                                                                                                                                                                                                                                                                          89.5%;
71.4%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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1.7e+06;
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cadherin
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RESULT 9
ABG33862
ID ABG3
XX ABG3
XX ABG3
XX ABG3
XX ABG3
XX BBG9
XX B Ly
XX B Syno

15-JUL-2002

(first entry)

ABG33862 standard; peptide;

B Lymphocyte Stimulator protein; B Lymphocyte Stimulator binding BLyS; biological fluid; serum; plasma; lymph; blood; urine; spina synovial fluid; saliva; mucus.

urine; spinal

Lymphocyte Stimulator (BLyS) binding peptide #436.

á 밁

CXPXTGC

Best Loc Matches Query Match

Local

Similarity 7; Conser

89.5%; Score 34; ilarity 100.0%; Pred. No. Conservative 0; Mismatc

Mismatches

0

Gaps

0

DB 5; 5, 1.7e+06; 0;

Length 7;

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amelioration of a disease or disorder associated with: aberrant B lymphocyte stimulator (BLyS), BLyS receptor expression or activity; cells of haematopoietic origin; or proliferative disease; and reducing, inhibiting or stimulating immunoglobulin production, B cell proliferation and graft rejection involving administration of BLyS binding polypeptides. The BLyS binding polypeptides are used in the treatment, prevention or amelioration of diseases such as immune system diseases, proliferative diseases, diseases of cells of hematopoietic origin, graft rejection, allergies, infectious diseases, arteriosclerosis, inflammatory disorders, hypergammaglobulinaemia, blood clotting disorders, ischaemia, and neurodegenerative diseases. The present sequence is a B lymphocyte stimulator protein binding peptide
                                                                                                                                                                                                                                                                                                             The treatment of various diseases e.g. rheumatoid arthritis, comprises administering B Lymphocyte stimulator binding polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunosuppressive; nephrotropic; antirheumatic; antiarthritic; neuroprotective; cytostatic; immunostimulant; anticumour; anti-HIV; antiasthmatic; antiallergic; thyromimetic; antianaemic; haemostatic; dermatological; antiinflammatory; cardiant; ophthalmological; uropathic; antidiabetic; antithyroid; antidepressant; hepatotropic.
Sequence
                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-AUG-2000; 2000US-0226700P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-AUG-2001; 2001WO-US025850
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ischaemia; graft-versus-host disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                          (HUMA-)
                                                                                                                                                                                                                                                                                                                                                                      2002-499775/53
                                                                                                                                                                                                                                                                              69; Page 233; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                       JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                          HUMAN GENOME SCI
 8
                                                                                                                                                                                                                                                                                                                                                                                                         Potter DM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= Pro, Tyr
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                                                                                                                                                                                                                                                                                                                                                                                                       Fleming
                                                                                                                                                                                                                                                                                                                                                                                                         Ţ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neurodegenerative disease;
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Synthetic.

WO200216412-A2

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RESULT 10
ABJ00545
ID ABJ005
XX ABJ000
XX ABJ000
XX O5-SE
XX D5-SE
XX B lym
XX B lym
XX B lym
XX antia
XW imchae
XW 
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a B Lymphocyte Stimulator (BLyS) binding polypeptide. BLyS binding peptides bind BLyS or BLyS-like proteins reversibly or irreversibly. The binding peptides are used in detection, isolation and/or purification of BLyS in a solution such as water or a buffer solution, as well as any fluid and/or cell obtained from an individual biological fluid, body tissue, body cell, cell line, tissue culture or other source containing BLyS or BLyS-like polypeptides. The biological fluid, solutions, lymph, blood, blood fraction, urine, synovisi fluid, spinal fluid, saliva and mucous. Sequences ABG33405-33415, ABG33423-33575, ABG33588-33846, ABG33848-33850 and ABG33852-33862 represent BLyS binding peptides of the invention
                                                                                                                                                                                                                                                            B lymphocyte stimulator protein binding protein; BLyS; immune disease; allergy; proliferative disease; infectious disease; arteriosclerosis; inflammatory disorder; hypergammaglobulinaemia; blood clotting; ischaemia; graft-versus-host disease; neurodegenerative disease; immunosuppressive; nephrotropic; antirheumatic; antiarthritic; neuroprotective; cycostatic; immunostimulant; antitumour; anti-HIV; antiasthmatic; antiallergic; thyromametic; antianaemic; haemostatic; dermatological; antiinflammatory; cardiant; ophthalmological; uropathic; antidiabetic; antithyroid, antidepressant; hepatotropic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-AUG-2000; 2000US-0226489P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-351647/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABJ00545 standard; peptide;
   Misc-difference
                                                           Misc-difference
                                                                                                                    Misc-difference
                                                                                                                                                                                                            Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CXPXTGC 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 stimulator protein binding protein consensus peptide #1.
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                                                                                                                                                Location/Qualifiers
                               /label= Ala, Glu, Met, Ser, Val
                                                                                        label- Ala, Asn, Lys,
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100.0%; Pred. No. 1.
ive 0; Mismatches
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hes 0;
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8
                                     Query Match
Best Local &
                            Matches
                                                                                            amelioration of a disease or disorder associated with: aberrant B lymphocyte stimulator (BLyS), BLyS receptor expression or activity; cells of haematopoietic origin; or proliferative disease; and reducing, inhibiting or stimulating immunoglobulin production, B cell proliferation and graft rejection involving administration of BLyS binding polypeptide. The BLyS binding polypeptides are used in the treatment, prevention or amelioration of diseases such as immune system diseases, proliferative diseases, diseases of cells of hematopoietic origin, graft rejection, allergies, infectious diseases, arteriosclerosis, inflammatory disorders, hypergammaglobulinaemia, blood clotting disorders, ischaemia, and neurodegenerative diseases. The present sequence is a conserved region of a B lymphocyte stimulator protein binding peptide
                                                                                                                                                                                                                                      The present invention relates to the treatment, prevention or amelioration of a disease or disorder associated with: aberra
                                                                                                                                                                                                                                                                         Claim 69; Page 230; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                             Beltzer JP,
                                                                                                                                                                                                                                                                                                                                                                                                              18-AUG-2000; 2000US-0226700P
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                                                                          Sequence
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                                                                                                                                                                                                                                                                                                             The treatment
                                                                                                                                                                                                                                                                                                                                                                                     (HUMA-) HUMAN GENOME SCI
                           Similarity
7; Conserv
  CXPXTGC
                                                                            13
                                                                            ă,
                            Conservative
                                                                                                                                                                                                                                                                                                                                                              Potter DM,
                                                                                                                                                                                                                                                                                                  of various diseases e.g. rheumatoid arthritis, comprises B Lymphocyte stimulator binding polypeptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= Ala,
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                                       100.0%;
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                                       Score 34;
Pred. No.
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                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gly,
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                                                                                                                                                                                                                                                                                                                                                               Rosen
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                            0,
                                                   Length 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ser
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                            Gaps
                             0,
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RESULT 11
ABG33861
ID ABG33861
AC ABG33
XX ABG33
XX ABG33
XX BG33
XX ABG33

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B Lymphocyte Stimulator protein; B Lymphocyte Stimulator binding peptide; BLyS; biological fluid; serum; plasma; lymph; blood; urine; spinal fluid; synovial fluid; saliva; mucus.

15-JUL-2002

entry)

B Lymphocyte

Stimulator (first

(BLyS)

binding peptide

WO200216412-A2 Synthetic

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ARESULT 12
ARU44932
ARU44932
ARU44932
ARU44932
ARU ARU44
AXX
AXX
ARU4932
ARU49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-AUG-2001; 2001WO-US025891.
                                                                                                                                                                      21-APR-2000;
02-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                      Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Propionibacterium acnes immunogenic protein #5828.
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     Skeiky YAW, I
L'maisonneuve
                                                                                                                                               07-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU44932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU44932 standard; protein; 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Page 121;
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                                                                                       (CORI-) CORIXA CORP
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7; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CXPXTGC 10
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  Persing DH,
re J, Zhang Y,
                                                                                                                                         2000US-0199047P.
2000US-0208841P.
2000US-0216747P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              osteopathic; neuroprotectant
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100.0%; Pred. No. 19
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fleming TJ,
Mitcham JL, Wang SS, Jen S, Carter D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central convous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (ELISA). Note: The sequence data for this patent did not form part of the printed specification, but was constant in the sequence data for obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                Mitcham JL, Sauthern Jen S, Jen S, Zhang Y, Wang S, Jen S, Zhang Y, Wallieve-Douglass J;
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N-PSDB; AAS59524.
                                                                                                                                                                                                                                                                                                                                         11-OCT-2002; 2002WO-US032727.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Acne vulgaris; antiseborrhoeic; dermatological;
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                                                                                                                                                                                                                                                                           15-OCT-2001; 2001US-00978825
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                                                                                                                                                                                                                     (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunostimulant; immune response; vaccine.
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Pred. No. 60;
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Benson DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC encoding a Propionibacterium acnes protein. The invention relates to an isolated polynucleotide (ACF64435-ACF64733) CC encoding a Propionibacterium acnes protein. The invention also relates to CC immunogenic fragments of P. acnes polypeptides. The invention CC immunogenic fragments of P. acnes polypeptides. The invention CC additionally encompasses expression vectors and host cells comprising a CC polynucleotide of the invention; antibodies against polypeptides of the invention; a ccine comprising a polypeptide of the invention; a ccine polypeptide and an isolated T cell population comprising T cells prepared CC via this method; a vaccine composition (comprising T cells prepared CC polynucleotides, antibodies, fusion proteins, T cell populations, or CC antigen-presenting cells that express the polypeptide); a method and kit CC for detecting or determining the presence or absence of P. acnes in a CC patient; and a method for inhibiting the development of P. acnes in a CC patient; The P. acnes solypeptides, polypeptides, polypeptides are useful for diagnosing, preventing or treating acne CC volgaris, T cell populations or antigen-presenting cells that express the polypeptides are useful for diagnosing, preventing or treating acne CC volgaris, or for stimulating an immune response specific for a P. acnes for cuclaic acid hybridisation. The vaccine composition is useful for the ethination of an immune response against P. acnes, or for treating acne, cc stimulation of an immune response against P. acnes, or for primers for cand the kit is useful for performing a diagnostic assay. The present CC proteins of the present a polypeptide predicted to be encoded by an ORF (open creading frame) contained within the P. acnes polypucleotides of the contained within the P. acnes polypucleotides of the format of the contained within the p. acnes polypucleotides of the 
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                                                                                                                                                                                                                                                                                            Cytostatic; Anti-inflammatory; Osteopathic; Neuroprotective; Nootropic; Gene Therapy; human; secretory protein; membrane proteins; cancer; inflammatory disease; osteoporosis; neurological disease.
                                                                                                                                                                                                                                                                                                                                                                             Human protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA54961 standard; protein; 113 AA.
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  (HELI-)
(REAS-)
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                                                             14-SEP-2001;
24-JAN-2002;
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                                                             2001JP-00328381.
2002US-0350435P.
                                                                                                                             2002EP-00006586
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  BIOTECHNOLOGY
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71.4%;
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Pred. No. 60;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; with vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunosupressave; antidiabetic; immunosupressave; antidiabetic; whypotensive; dermatological; immunosuppressive; antidiabetic; whypotensive; dermatological; immunosuppressive; antidiabetic; whypotensive; dermatological; immunosuppressive; antidiabetic; white interactive; antidiabetic; continued; antidiated; antidiated; antidiated; antidiated; whypotension; while interactive disorder; hypotension; while interactive disorder; osteoarthritis; graft vs host disease; when encodegenerative disorder; osteoarthritis; graft vs host disease; when conditions disease; diabetes mellitus; hypothyroidism; SCID; AIDS; while immunodeficiency; malaria; autoimmune disorder; asthma; where combined immunodeficiency; malaria; autoimmune disorder; asthma; where combined immunodeficiency; malaria; autoimmune disorder; asthma; where combined immunodeficiency; malaria; autoimmune disorder; asthma; whose damage; cartilage damage; antiinflammatory disease; coagulation; where the cartilage damage; antiinflammatory disease; coagulation; where the combined is contraceptive.
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Matches 5
                                                                                                                                                                                   31-MAR-1999;
02-APR-1999;
05-APR-1999;
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Yamamoto
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seki N,
                                                                                                                                                                                                                                                                                                                                                                      05-OCT-2000.
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                                                                                               (CURA-) CURAGEN CORP
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Hio Y, Otsuka K, Nagai
Otsuka M, Nagahari K, I
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R,

Leach M;

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Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease.
                                                                                                                                                                                                    WPI; 2000-602362/57.
N-PSDB; AAC76236.
                                                                                                                                                            Claim 11, Page 2743; 5507pp; English.
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CC which represent the human ORFX open reading frames 1 to 3161. The ORFX CC sequences have activities such as: cytostatic; hepatotropic; vulnerary; CC antipsoriatic; antiparkinsonian; nootropic; neuroprotective; osteopathic; CC anticonvulsant; antiarthritic; immunosuppressant; immunostimulant; CC cardiant; thrombolytic; coagulant; vascotropic; antidiabetic; hypotensive; CC dermatological; immunosuppressive; antiinflammatory; antidiabetic; The cc antiviral; antifungal; antitheumatic; antithyroid; and antianaemic. The sequences can be used for determining the presence of or predisposition to, or preventing or treating pathological conditions associated with an CC ORFX-associated disorder. The nucleic acids can be used to express ORFX CC proteins in gene therapy vectors. The proteins and nucleic acids may be used to treat cancers, proliferative disorders, neurodegenerative considers, osteoarthritis, graft vs host disease, cardiovasqualar disease, estorage, systemic lupus erythematosus, severe combined immunodeficiency consorders, asthma, allergies, aplastic anaemia, burns, wounds, bone and cartilage damage, nocturnal haemoglobinuria, antiinflammatory disease; to enhance coagulation; to inhibit thrombosis; and as a contraceptive

Sequence 138 AA;

Query Match 89.5%; Best Local Similarity 71.4%; Matches 5; Conservative 1 CXPXTGC 7 0; Mismatches Score 34; DB 3; Length 138; Pred. No. 1.3e+02; O; Mismatches 2; Indels 0 Gaps 0,

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